

# SEQUENCE LISTING

<110> Quinn, Kerry E.  
Pena, Carol A. E.  
Li, Li  
Spaderna, Steven K.  
Leite, Marlo W.

<120> Aortic Carboxypeptidase-Like Proteins and Nucleic Acids  
Encoding Same

<130> 15966-581 CIP

<140> 09/996,015

<141> 2001-11-28

<150> 09/641,741

<151> 2000-08-18

<150> 60/159,613

<151> 1999-10-14

<150> 60/175,534

<151> 2000-01-11

<150> 60/224,086

<151> 2000-08-09

<160> 47

<170> PatentIn Ver. 2.1

<210> 1

<211> 2382

<212> DNA

<213> Homo sapiens

<400> 1

```
gcggggggca ggaaggggcg gggggctcgg cgcactcggc aggaagagac cgaccogcca 60
ccgcgcgtag ccgcgcgcgc cctgggaact aatccccgcc atgtgggggc tcctgctcgc 120
cctggcgcgc ttgcgcgcgc ccgtcggccc ggctctgggg gcgcgcagga actcgggtgt 180
gggcctcgcg cagcccgga ccaccaaggt ccaggctcg accccggccc tgcatagcag 240
ccgcgcacag ccgcgcgcgc agacagctaa cgggacctca gaacagcatg tcgggattcg 300
agtcatacag aagaaaaagg tcattatgaa gaagcggaag aagctaacct taactcgccc 360
caccacactg gtgaactgcg ggccccttgt gacccccact ccagcagga ccccgagccc 420
cgctgagaaa caagaaacag gctgtcctcc tttgggtctg gactccctgc gactttcaga 480
tagccggctt gaggcataca gcagccagtc ctttggtctt ggacacacc gaggaocgct 540
caacattcag tcaggcctgg aggacggcga tctatatgat ggagcctggt gtgctgagga 600
```

```

gcaggacgccc gatccatggt ttcaggtgga cgttgggcac cccacccgct tctcgggtgt 660
tatcacacag ggcaggaact ctgtctggag gtatgactgg gtcacatcat acaagggtcca 720
gttcagcaat gacagtcgga cctgggtgggg aagtaggaac cacagcagtg ggtatggacgc 780
agtatttctt gccaattcag acccagaaac tccagtgtctg aacctctctg cggagccccca 840
ggtggccccc ttcattcgcc tgggtcccca gacctggctc cagggaggcg cgccttgccct 900
ccgggcagag atcctggcct gccagttctc agaccccaat gacctattcc ttgaggcccc 960
tgcgtcggga tctctgacc ctctagactt tcagcatcac aattacaagg ccattgaggaa 1020
gctgatgaag caggtacaag agcaatgccc caacatcac ccgcatctaca gcattgggaa 1080
gagctaccag ggccctgaagc tgtatgtgat ggaaatgtcg gacaagcctg gggagcatga 1140
gctgggggag cctgaggtgc gctacgtggc tggcatgcat gggaacgagg ccttggggcg 1200
ggagttgctt ctgctctctg tgcagttctt gtgccatgag ttcttgcgag ggaacccacg 1260
ggtgaccocg ctgctctctg agatgcgcct tcacctgctg cctcccatga accctgatgg 1320
ctatgagatc gctaccacc ggggttcaga gctgggtggc tgggcccagg gccgctggaa 1380
caaccagagc atcgatctta accataattt tgcagacctc aacacaccac tgtgggaagc 1440
acaggacgat gggaagggtc cccacatcgt ccccaaccat cacctgccat tgccactta 1500
ctacaccctg cccaatgcca ccgtggctcc tgaaacgcgg gcagtaatca agtggatgaa 1560
gcggatcccc tttgtgctaa gtgccaacct ccacgggggt gagctcgtgg tgcctaccc 1620
attcgacatg actcgacccc cgtgggctgc ccgcgagctc acgcccacac cagatgatgc 1680
tgtgtttcgc tggctcagca ctgtctatgc tggcagtaat ctggccatgc aggacaccag 1740
ccgcgcagccc tgccacagcc aggacttctc cgtgcacggc aacatcatca accgggctga 1800
ctggcacacg gtccccggga gcatgaatga cttcagctac ctacacacca actgctttga 1860
ggtcactgtg gagctgtcct gtgacaagtt cctcacgag aatgaattgc cccaggagtg 1920
ggagaacaac aaagaagccc tctcaccta cctggagcag gtgcgcatgg gcattgcagg 1980
agtgttgagg gacaaggaca cggagcttgg gattgtgac gctgtcattg ccgtggatgg 2040
gattaaccat gacgtgacca cggcgtgggg cggggattat tggcgtctgc tgaccccagg 2100
ggactacatg gtgactgcca gtgccgaggg ctaccattca gtgacacgga actgtcgggt 2160
cacctttgaa gagggccctt tccctgcaa tttcgtgctc accaagactc ccaaacagag 2220
gctgcgcgag ctgctggcag ctggggccaa ggtgcccccg gaccttcgca ggcgcctgga 2280
gcggctaagg ggacagaagg attgatacct gcggtttaag agccctaggg caggctggac 2340
ctgtcaagac gggaagggga agagtagaga gggagggaca aa 2382

```

<210> 2

<211> 734

<212> PRT

<213> Homo sapiens

<400> 2

```

Met Trp Gly Leu Leu Leu Ala Leu Ala Ala Phe Ala Pro Ala Val Gly
  1                      5                      10                      15

```

```

Pro Ala Leu Gly Ala Pro Arg Asn Ser Val Leu Gly Leu Ala Gln Pro
      20                      25                      30

```

```

Gly Thr Thr Lys Val Pro Gly Ser Thr Pro Ala Leu His Ser Ser Pro
    35                      40                      45

```

```

Ala Gln Pro Pro Ala Glu Thr Ala Asn Gly Thr Ser Glu Gln His Val

```

50	55	60
Arg Ile Arg Val Ile Lys Lys Lys Lys Val Ile Met Lys Lys Arg Lys		
65	70	75 80
Lys Leu Thr Leu Thr Arg Pro Thr Pro Leu Val Thr Ala Gly Pro Leu		
	85	90 95
Val Thr Pro Thr Pro Ala Gly Thr Leu Asp Pro Ala Glu Lys Gln Glu		
	100	105 110
Thr Gly Cys Pro Pro Leu Gly Leu Glu Ser Leu Arg Val Ser Asp Ser		
	115	120 125
Arg Leu Glu Ala Ser Ser Ser Gln Ser Phe Gly Leu Gly Pro His Arg		
	130	135 140
Gly Arg Leu Asn Ile Gln Ser Gly Leu Glu Asp Gly Asp Leu Tyr Asp		
	145	150 155 160
Gly Ala Trp Cys Ala Glu Glu Gln Asp Ala Asp Pro Trp Phe Gln Val		
	165	170 175
Asp Ala Gly His Pro Thr Arg Phe Ser Gly Val Ile Thr Gln Gly Arg		
	180	185 190
Asn Ser Val Trp Arg Tyr Asp Trp Val Thr Ser Tyr Lys Val Gln Phe		
	195	200 205
Ser Asn Asp Ser Arg Thr Trp Trp Gly Ser Arg Asn His Ser Ser Gly		
	210	215 220
Met Asp Ala Val Phe Pro Ala Asn Ser Asp Pro Glu Thr Pro Val Leu		
	225	230 235 240
Asn Leu Leu Pro Glu Pro Gln Val Ala Arg Phe Ile Arg Leu Leu Pro		
	245	250 255
Gln Thr Trp Leu Gln Gly Gly Ala Pro Cys Leu Arg Ala Glu Ile Leu		
	260	265 270
Ala Cys Pro Val Ser Asp Pro Asn Asp Leu Phe Leu Glu Ala Pro Ala		
	275	280 285
Ser Gly Ser Ser Asp Pro Leu Asp Phe Gln His His Asn Tyr Lys Ala		
	290	295 300
Met Arg Lys Leu Met Lys Gln Val Gln Glu Gln Cys Pro Asn Ile Thr		

305		310		315		320
Arg Ile Tyr Ser Ile Gly Lys Ser Tyr Gln Gly Leu Lys Leu Tyr Val						
	325		330		335	
Met Glu Met Ser Asp Lys Pro Gly Glu His Glu Leu Gly Glu Pro Glu						
	340		345		350	
Val Arg Tyr Val Ala Gly Met His Gly Asn Glu Ala Leu Gly Arg Glu						
	355		360		365	
Leu Leu Leu Leu Leu Met Gln Phe Leu Cys His Glu Phe Leu Arg Gly						
	370		375		380	
Asn Pro Arg Val Thr Arg Leu Leu Ser Glu Met Arg Ile His Leu Leu						
385		390		395		400
Pro Ser Met Asn Pro Asp Gly Tyr Glu Ile Ala Tyr His Arg Gly Ser						
	405		410		415	
Glu Leu Val Gly Trp Ala Glu Gly Arg Trp Asn Asn Gln Ser Ile Asp						
	420		425		430	
Leu Asn His Asn Phe Ala Asp Leu Asn Thr Pro Leu Trp Glu Ala Gln						
	435		440		445	
Asp Asp Gly Lys Val Pro His Ile Val Pro Asn His His Leu Pro Leu						
	450		455		460	
Pro Thr Tyr Tyr Thr Leu Pro Asn Ala Thr Val Ala Pro Glu Thr Arg						
465		470		475		480
Ala Val Ile Lys Trp Met Lys Arg Ile Pro Phe Val Leu Ser Ala Asn						
	485		490		495	
Leu His Gly Gly Glu Leu Val Val Ser Tyr Pro Phe Asp Met Thr Arg						
	500		505		510	
Thr Pro Trp Ala Ala Arg Glu Leu Thr Pro Thr Pro Asp Asp Ala Val						
	515		520		525	
Phe Arg Trp Leu Ser Thr Val Tyr Ala Gly Ser Asn Leu Ala Met Gln						
	530		535		540	
Asp Thr Ser Arg Arg Pro Cys His Ser Gln Asp Phe Ser Val His Gly						
545		550		555		560
Asn Ile Ile Asn Gly Ala Asp Trp His Thr Val Pro Gly Ser Met Asn						

565	570	575
Asp Phe Ser Tyr Leu His Thr Asn Cys Phe Glu Val Thr Val Glu Leu		
580	585	590
Ser Cys Asp Lys Phe Pro His Glu Asn Glu Leu Pro Gln Glu Trp Glu		
595	600	605
Asn Asn Lys Asp Ala Leu Leu Thr Tyr Leu Glu Gln Val Arg Met Gly		
610	615	620
Ile Ala Gly Val Val Arg Asp Lys Asp Thr Glu Leu Gly Ile Ala Asp		
625	630	635
Ala Val Ile Ala Val Asp Gly Ile Asn His Asp Val Thr Thr Ala Trp		
645	650	655
Gly Gly Asp Tyr Trp Arg Leu Leu Thr Pro Gly Asp Tyr Met Val Thr		
660	665	670
Ala Ser Ala Glu Gly Tyr His Ser Val Thr Arg Asn Cys Arg Val Thr		
675	680	685
Phe Glu Glu Gly Pro Phe Pro Cys Asn Phe Val Leu Thr Lys Thr Pro		
690	695	700
Lys Gln Arg Leu Arg Glu Leu Leu Ala Ala Gly Ala Lys Val Pro Pro		
705	710	715
Asp Leu Arg Arg Arg Leu Glu Arg Leu Arg Gly Gln Lys Asp		
725	730	

<210> 3  
 <211> 20190  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (2206)  
 <223> Wherein n is A, C, G, or T

<230>  
 <231> misc\_feature  
 <232> (2207)  
 <233> Wherein n is A, C, G, or T

<220>  
<221> misc\_feature  
<222> (2209)  
<223> Wherein n is A, C, G, or T

<220>  
<221> misc\_feature  
<222> (2211)  
<223> Wherein n is A, C, G, or T

<220>  
<221> misc\_feature  
<222> (2214)  
<223> Wherein n is A, C, G, or T

<220>  
<221> misc\_feature  
<222> (2215)  
<223> Wherein n is A, C, G, or T

<220>  
<221> misc\_feature  
<222> (2218)  
<223> Wherein n is A, C, G, or T

<220>  
<221> misc\_feature  
<222> (2220)  
<223> Wherein n is A, C, G, or T

<220>  
<221> misc\_feature  
<222> (2221)  
<223> Wherein n is A, C, G, or T

<220>  
<221> misc\_feature  
<222> (2222)  
<223> Wherein n is A, C, G, or T

<220>  
<221> misc\_feature  
<222> (2224)  
<223> Wherein n is A, C, G, or T

<220>  
<221> misc\_feature  
<222> (2226)

<223> Wherein n is A, C, G, or T

<220>

<221> misc\_feature

<222> (2227)

<223> Wherein n is A, C, G, or T

<220>

<221> misc\_feature

<222> (2229)

<223> Wherein n is A, C, G, or T

<220>

<221> misc\_feature

<222> (2230)

<223> Wherein n is A, C, G, or T

<400> 3

```
atgtgggggc tectgtctgc cctggcgcgc ttgcgcgcgc cgtcgggccc ggctctgggg 60
ggcgcacagga actcgggtgt gggcctcgcg cagcccggga ccaccaaggt cccaggctcg 120
accccgggccc tgcatagcag cccggcacag ccgcggcgag agacagctaa cgggacctca 180
gaacagcatg tccggattcg agtcatcaag aagaaaaagg tcattatgaa gaagcggaag 240
aagctaactc taactcgccc caccocactg gtgactgccc ggccccttgt gacccocact 300
ccagcaggga ccctcgaccc cgtcgagaaa caagaaacag gctgtcctcc tttgggtctg 360
gagtcctctg gagtttcaga tagccggctt gaggcattca gcagccagtc ctttgggtctt 420
ggaccacacc gaggaagggt caacattcag tcaggcctgg aggacggcga tctatatgat 480
ggagcctggt gtgctgagga gcaggacgcc gatccatggt ttcagggtgga cgtggggcac 540
cccacccgct tctcgggtgt tatcacacag ggcaggaaact ctgtctggag gtatgactgg 600
gtcacatcat acaagggtcca gttcagcaat gacagtcgga cctgggtgggg aagtaggaac 660
cacagcagtg ggatggagcg agtatttctt gccaatcag acccagaaac tccagtgtctg 720
aacctcctgc cggagcccca ggtggcgcgc ttcattcgcc tgcctgcccc gacctggctc 780
cagggaggcg cgccttgccct ccgggcagag atcctggcct gccagtcctc agaccccaat 840
gacctattcc ttgaggcccc tgcgtcgga ttctctgacc ctctagactt tcagcatcac 900
aattacaagg ccattgaggaa gctgatgaag cagggtacaag agcaatgccc caacatcacc 960
cgcattctaca gcattgggaa gagctaacag ggcctgaagc tgtatgtgat ggaaatgtcg 1020
gacaagcctg gggagcatga gctgggggag cctgagggtgc gctaagtggc tggcatgcat 1080
gggaacgagg ccctggggcg ggagttgctt ctgctcctga tgcagttcct gtgccatgag 1140
ttcctgcgag ggaacccacg ggtgacccgg ctgctctctg agatgcgcat tcacctgctg 1200
ccctccatga accctgatgg ctatgagatc gcctaccacc ggggttcaga gctgggtggg 1260
tgggcgcgag gccgctggaa caaccagagc atcgatctta accataattt tgcctgacct 1320
aacacaccac tgtgggaagc acaggacgat ggggaagggtgc cccacatcgt ccccaacct 1380
cacctgccat tgcccactta ctacacccctg cccaatgcca ccgtggctcc tgaaacgcgg 1440
gcagtaatca agtggatgaa gcggatcccc tttgtgctaa gtgccaacct ccacgggggt 1500
gagctcgtgg tgtcctaccc attcgacatg actcgacccc cgtgggctgc ccgcgagctc 1560
acgcccacac cagatgatgc tgtgtttcgc tggctcagca ctgtctatgc tggcagtaat 1620
ctggccatgc aggacaccag ccgcgcaccc tgcacagcc aggaattctc cgtgcacggc 1680
aacatcatca acggggctga ctggcacacg gtcccgggga gcatgaatga cttcagctac 1740
ctacacacca actgctttga ggtaactgtg gagctgtcct gtgacaagtt ccctcacgag 1800
```

aatgaattgc cccaggagtg ggagaacaac aaagacgccc tcttcaccta cctggagcag 1860  
gtgcgcattg gcaattgcagg agtgggtgagg gacaaggaca cggagcttgg gattgctgac 1920  
gctgtcattg ccgtggatgg gattaacat gacgtgacca cggcgtgggg cggggattat 1980  
tggcgtctgc tgaccccagg ggactacatg gtgactgcca gtgcggaggg ctaccattca 2040  
gtgacacgga actgtcgggt cacctttgaa gagggcccct tcccctgcaa ttctgtgctc 2100  
accaagactc ccaaacagag gctgcgcgag ctgctggcag ctggggccaa ggtgcccccg 2160  
gaccttcgca ggcgcctgga gcggctaagg ggacagaagg attgannant ncannttnan 2220  
nntngnnann tctcacttat aaatggaagc tggcgggaca cgggtggctca ctctgtaat 2280  
cccaacactt tgggaggctg aggcgggtgg atcacgaggt caggagatcg agaccatcct 2340  
gactaacaag gtgaaacccg tctctactaa aaacacaaaa aattagctgg gcgtggtggc 2400  
ggcacctgta gteccagcta ctcgggaggc tgaggcagga gaatggcatg aaccagagg 2460  
tcggagcttg cagtgcgcg agttcacgcc actgcattcc agcctgggca acagagcag 2520  
actctgtctc aaaaaaata aattaaataa aaataaataa atggaaacta agctgtgggt 2580  
atgcaaaggc atacagaatg gtataatgga cattggagac tcagaaggag gagggtaagc 2640  
gggggggtgac agataaaaaa aactgcatgt tgcatacaat gtacactact cgggtgatgg 2700  
gcgctctaag atttcaaaact tcaccactat acagttctcc cctgtaacca aaaaccgctg 2760  
gtacccctaa agcaattgaa ataaaaatag aaactatggt gtagcctgga tgacatagcg 2820  
aaaacttgct tottaaaaaa aaaaaaatgt ggccgggtgc agtggctcac acctgtaatc 2880  
ccagcacttt gggaggccca aggcgggcag atcacaaggt caggagattg agaccgtcct 2940  
ggctaacaag gtgaaactcc atctctacta aaaatacaaa aaattagccg ggtgtggtgg 3000  
cacacgcctg taatccagc tacttgggag gctgaggcag gagaatcgct tgaaccaga 3060  
ggcggagggt gcagtgcgc gagatcgac cacagcactc cagcctggtg acagagtgcg 3120  
atttagtctc aaaaaaaaaa aaaaaaaaaa aaaaaaaag gtagaaatta gctgagcgtg 3180  
gtgacacgct ccagataactt gggaggctga ggtgggagga tcgcttgaa cccaggagt 3240  
cagactgcag tgagctgtga ttacactatt gcactccagc ctaggctgtg ggaaagagag 3300  
tttctggggg gccagctgag ttagtcttcc ctgtgtgaga caccatggg aagccatgcg 3360  
cggcctctga ggagaaaagt ctcttattg ccttcatgtc tttaacgccg agagcagaac 3420  
ccctcagcgg cattccacag gttgctcagg catataacac tcccttgaa cagtggagta 3480  
taatcaaaaca tcttggctcc tctgaaacc cactcccacc cgtttcagtc ccgataagtt 3540  
aaagatttgt tttgttttgt tttgtttga gacggagtct cgctctgtcg cccaggctgg 3600  
agtgcgggtg ctgatctcg gctcactgca agctccgct cccgggttca cgccattctc 3660  
ccgcctcagc ctcccgagta gctgggacga caggcgcccg ccaccacgcc cggctaattt 3720  
tttgtatttt tagtagagac ggggtttcac cacgttggcc agagtggctc cgaactcctg 3780  
acctcaagcg atccaccac ctccgcctcc caagtgtcg ggattacagg cgtgagccac 3840  
cgcgcccggc cagttaaaga tcttaagtag ttgacactc ctctttgctc aaggaaattc 3900  
acagaaaccg ccaactgctat acatcttaca gaatgactct ccagttctcc ttactgatt 3960  
aatcctttcc ctcatccctt cctcctctc ccatctgcc taagaacaaa gagcttgtaa 4020  
accaataaat tgggcggagc ctgagaactc tgggcggtga gcaagcctcc gacgctccgg 4080  
tcccctggac ccgcctttta aacgcttatt ctgtctcttt ctaactcctt tgtctccgcc 4140  
ggactcgggg taaccgctag gcgttatggg gctgttttcc ccaacatagg caacagagca 4200  
ggacagtgtc tctaaaaaaa caaaacaaaa actatatatt gtactattct gataaaaatg 4260  
acttagttac aaacaaagaa caaatcaaca gatagtcatg ctgtggagat caggaatatt 4320  
ccttcccagg gtaaatgaaa gaccaattcc ctaacgtcat gtggatatac gcttgtggct 4380  
taagataaaa ttaccgctga cagcatcaaa taccagggat aaaactcagt cttcaacacg 4440  
catatgtatc tctgggggtt gaatcctctg gaggtcttgt taaaaatgca gattctggtc 4500  
aagagttcga gaccagcctg gccaatatgg tgaaacctg tctctactaa aaacacaaaa 4560  
attagctggg tgtgggtgtg gacgcctgta gtcccagcta ctcaggagac tgaggcagga 4620  
gaattgcttg aaccggggag gtggcagttt agtgagctga gatcgggcca ctgcactcca 4680



gcctgggaga	cagagtgaga	ctctgtcaaa	aaaaaaaaaa	aaaaaaaaatg	catattctga	4740
ttcaataggt	ctggggcaga	ggtgtttttt	ttgtttgttt	gttttttgtt	ttttggtttt	4800
ttttttgggt	tttttttttt	gacagagtct	agctctttca	cctaggctgg	agtgcattgac	4860
accatcccag	ctcactgcaa	cctccgcttc	ttgggttcaa	gogattctcc	tgcctcagcc	4920
tctgaatag	ctgggattac	aggcgtgcac	caccaccccc	agctaagttt	tgtatttgta	4980
gtagagatgg	ggtttcacccg	tgttgccag	gtaagttttt	tatttgatt	tggctctgaa	5040
ctcctgacct	caggatgatcc	gcccgcctcg	gcctcccaaa	gttctgggat	tacaggcgtg	5100
agccactgca	cccgccctgt	tctgcatttc	taacaagttt	ccaggggatg	ctactgctgc	5160
tggctctcaa	ccacactttg	tggagcaagg	ctctcaaaga	ccttgatgta	tgtaggagag	5220
aaagctgggg	tagagagtga	tgaggggaga	acgggtgcgt	gggagatgc	tcccctgtgc	5280
atcctggtcc	catgtgaggc	tccaacaatg	ctcacctaca	tcacagggag	agcacctagc	5340
aggaaatgag	ttctgcttta	gcattccaggc	acaggagatt	agaggcacag	gcaggcagta	5400
gattctactt	cattatttgt	gcagctggac	acagagcttc	ctttcttttc	cttgatactg	5460
ttttattcca	tctaagtatg	taggagtaag	agggtgtgt	tacactgttt	tccccacctt	5520
taatgcatct	gatcaacctt	ggagccccct	aagacctat	attatctcac	tttatcatca	5580
cagcaaacct	gggagaagga	tatggttctt	gttttacaga	tgaggaaact	aagtctcagg	5640
gaggtgaaac	tactgcccaa	ggatagccaa	acaaaataca	cgtcagaagt	gggatgtgaa	5700
acgaagcctg	tatgtcacca	gagtcacctt	tctctcccc	ctccaaccac	ctaaccacac	5760
caggaggttg	gcaggagatt	cctagcccac	cccttacatt	aaaatccctt	ttaggcgggt	5820
gccactatcc	agtccttctc	aattgcacct	agtgcagcca	cgaaagatct	tctacctggc	5880
tcttggtaga	tgagatctgg	ctatacaggt	acttggtgct	aaacctgccc	ctctgcccct	5940
ggagctatca	cctccagatc	ctgctacttg	tacctttgca	gccccaggta	gccagtggca	6000
agggccaggg	gtggcagcag	ggctgggagt	ggagaagagt	gtgagaaagt	gctgcggggc	6060
tcaggagaca	cagcagggaa	ccaaggggtc	ctaaggggtg	caatagagga	caggggcagg	6120
gagtgcagag	tgggtgggaag	ggggatggga	gctgggtgca	ggagacataa	gagatggagc	6180
atcccgccca	cacacgggtg	ctcacacctg	ttatcccagc	actttgggag	gccgaggtgg	6240
gtggatcacg	aggtcaggag	atcgagacca	tcttggttaa	cacggtgaaa	ccccgtctct	6300
actaaaaaca	caaaaaatta	gccaggcgag	gtggtgtgca	cctgtagtcc	cagcttcttg	6360
ggaggctgag	gcaggagaat	ggcgtgaacc	caggaggtgg	agcttgacgt	cagctgagat	6420
cccgccactg	cattccagcc	tgggtgacag	agtgcagctc	gtctcaaaaa	aaaaaagaat	6480
aaaagaaaaa	agaggtggag	catcctgcag	ccctggcccc	taaaagattg	gtgggagagt	6540
gccagctgct	ccaccctagt	cactttggga	actggtcttt	cagttcacgg	cctgccatgt	6600
cctctcctgc	aaatcctggc	actgttgagg	aggctccttt	agccctgggt	tgtccactct	6660
aaccttgaat	atattataca	cacactttat	gagagctgac	gagggaccag	gtgctgttct	6720
aggctctgag	gtgcagctgt	ggacattttg	gtacaaagtt	cttctggcag	ggtacttacc	6780
tctgtctggg	ggtgggggaa	cctgaacagc	caacacataa	gtaaagcaag	atcatctcgg	6840
tgttgagtgc	cttgaagaca	ataatttaaa	cgggtgggag	gatagagtgt	gtgaagtga	6900
aaagtttgct	ttagtcaggg	tagtcaggga	aagccttttg	gagcaggtga	tattgaaagg	6960
aaatctgact	gagaaggcaa	attccatgca	caaattaaaa	ggccaggagg	ctagttgggc	7020
tgttgctggg	gaggagcagc	tagaatgccg	gagtgactgg	gggagatggg	gccaggggat	7080
agggaggcag	atggaatggg	aaaggcgtgg	gcaggaagaa	cttggtcatt	aagaccttgc	7140
aggtgaaccc	actggggcct	taagcctgga	ggaacttgac	agaatttgcc	tactgtgtgg	7200
ggaacggctt	ggaggggggtg	tgggcttcag	gaggtgaga	tgtcctgttt	cttgtgcccc	7260
ctcctttctt	cccaacaccc	gagaaacctg	gatgggtgtg	gggaccagag	acctggaggt	7320
ggccagattg	ggctttggcg	ggacgcttag	cagccctcgg	gacctgttca	gactgcggcc	7380
tcccaccttc	gggaagcatt	ggcgtgccc	atctgcccct	gcctggcgct	cagggagctc	7440
cggctgtgca	gcgtctccct	tgaatgtct	ctctgtcttc	ccatccagtg	cctgggaccc	7500
ggcagcgccg	tcagggcagg	gggctgcgag	gcgggaccca	gttgcacgtg	ggccctgtgg	7560

ggtaactccc tttaggggggt cctctagctc ttacccctgc gggggtgggg gagaccagat 7620  
 gcttcgagga gctccaggac cagtgcctat ggggtagtcc ctgggggggg tgggccccag 7680  
 tcccagactg cgggcgccta tttctttctg ggggttcgtgt gagggtgggc tgccagaatg 7740  
 gtgccacaaa gctgcttttg ggtgattcaa atcatttata cagatagtgc ccctgcaaaa 7800  
 aacatttgcg caggggccccg cttacgccag aggattggcg gccacttctg ggcatcgctc 7860  
 ctggtgggga tgggagcatt tccctggaga gccctttgca aaggccaagc gccggccaaa 7920  
 ggcacacgcg tggacgcgtt tccctccttc tggagagatg accaggaatg caggatccaa 7980  
 aggggggtctt ggagggaggg cgggaagggc atctccgat ctgggcagac ccagggtctg 8040  
 cggctccccg aggagaatac gggctggggg caggagaccg gagggcaggt caggcagtgc 8100  
 atcaaccctt ggctcctcca ccgcagcccc agcccgagc ctatcgctca ggcttctctc 8160  
 tccgggttat gtaaccccg gacgggacgt ggcagccggg tgagtgagcg aaggagttag 8220  
 ggaggggaagg gaaaggagag gaggggcagg gccgggcttg gtgatggtgg tgggtgggaag 8280  
 cgcgcgcgtg ccgcctcttc ttgggcccc tgggttgtct ttctggagga ttccgggacc 8340  
 agccctctcc ccaggtccg ggtcgcccc tagcccccg ccgcctcatt tcccttcac 8400  
 tcttttcccc cttctgtccc acccgccctg ccaggggggc tctggctctg gatagctttt 8460  
 cctctccggt tgtagtttcc ttcccaaagt tctcagcttt gctacctgc ccaagtcatt 8520  
 agccgtctg agcctcagtt tatcagtttg taaaatgaag ttgattgag cggccacgtg 8580  
 taaaactcct ggcatagtgc atggtacaaa gtagatgtct gctgcaggct aagggcctcg 8640  
 aggggctaag tgaatgttg tgtgccaggc tgggtgtcag agccccggga gccgcagcca 8700  
 cgaatggttg gctcccggtt ggtaaaagaa tttatcaaca acagtatagg ttgaaaagt 8760  
 tttattagat ggaaagaact ccacagcaga gcgcagcggg atgcttcggc aagagaggcc 8820  
 tgagctcact tgcagggaac tgaagggtaa ttttgaccac attagttttg taggtcatag 8880  
 taaatgatta catttgtaga cattttgga ccttgatgac agcaaagggt gcacaatggg 8940  
 ttccaacatg cgtgcattcc ggagatgtat agaaattcta gggaaagaag cctggtacca 9000  
 gatgtggctt tagataatag gaaagtacca ttctgagttc ttccagataag gtgctttgcc 9060  
 tctgatggt ctgcttgatg gccaccagggt gatccttgct ctccctcatt tccccctgat 9120  
 aaatatattg ggcaaatctt tgaccctttg tatttctcca tgcctcatgt tacttgtctg 9180  
 ttaggatccc aagaaaggga aaatggcaca gtgaagaggg gtgtccagtc tatctggcta 9240  
 ctctctgctg aaaaggggca ttgaaaggat tcccttcttg ctttctgtca tgaagggaat 9300  
 gaagggtcat gataaacttg ttcatggagg gaagaccaga ttccatcaag agggcccatg 9360  
 aaaatagaag ttgctgttgc aggtctggtat tgggattgca tagtcatctg taggtggaat 9420  
 cattgtaagc tggagatat aagcattaaa aggcagggaat taccggcatg cacctccatg 9480  
 cccacagatt tttgtgttt tagtagagac aggttctcac catgttggcc aggtctggtc 9540  
 ccaactcctg acctcaggtg atccgccccg ctgccttggt tctcccaaag tactaggatt 9600  
 acaggtgtga gcaaccacac ctggccccctg gggctctcaat ttgtgtatt atgcatggcc 9660  
 tccaccagtc tagcttgga aagggcaggg ctttcagata gtttcataca tacaaaatta 9720  
 ttatttcttt ttattttatt ttatttgaga tggaaatttg ctcttggtgc ccaggctgga 9780  
 gtgcagtggc gcaatctcag ctccaccaca cttccgctc caaggttcaa acgattctcc 9840  
 tgccctcagc tctggagtaa ctgggattac aggcattgac caccatgccc agctcatttt 9900  
 gtatttttag tagagatggg gtttctccgt gttggctagg ctggtctcaa acctcaggtg 9960  
 acccgctcgc ctccagctcc caaagtgtct ggattacagg tgtgagccac cgcgccagc 10020  
 tattatttct tataatttag aaaaattaac aggttttatt atatatattt cattccctcc 10080  
 aacagagaag ttaccatatg atcctgtctg ccttaacctc tgtttgggcc agaattggtg 10140  
 gcctggtatt gccaataggt tctatgttgg ggacagcttc tgcccagctc tgttatttag 10200  
 actgggagca tgagcttcat ctgcccattg tgaagatcac acgtgtgatt ttttgtgtgt 10260  
 gggaacagca ggtagttaat accacaaata catcttgcca ggttaaata aaggcaacag 10320  
 ttaaagtctg aaattcttga atgaacttag agggatcctg actaaatgaa cccaacttgg 10380  
 attgaatttg caaaagatca gacatgatca gaaaagggac atgaacttgg ctgttccca 10440

aatcttcatt	agccacotta	gggagaggca	aaatatattt	gggatttttc	tgaggactct	10500
gtactagtag	catatgtgac	tcccttgaga	gtatgtgaag	gggagaaagt	atttggttat	10560
gtgggtggga	gattgaactag	ggaatggagc	agatggagag	ggtgtagggt	aagagtgagc	10620
aggttgagga	ggatgtaata	ggcaaaagga	aggatcatct	aagacatcag	aaccgggaag	10680
ggaggacgtt	ccttggaagc	atacatgaca	atttgtatgt	aattttgggt	ttggatttgg	10740
ggataaagca	aaaaagacct	gaacatatgg	gactttctgaa	tcttttccaa	ggttccggca	10800
aaaaatcagt	taagttgtaa	agtagcattg	caatcccaag	tttcattaat	tggccaaatt	10860
gattgattag	ggagcttgta	ttgaacccaa	gcaatattag	aaaaaaggat	atgcttttta	10920
aactcttatt	tattttttat	ttgtattttt	tgagacagag	tcttgctgtg	tgcgccaggc	10980
tggagtgtct	tggcgccatc	ttggcccact	gcaacctccg	cccccggggt	acaagtgatt	11040
ctcctgcctc	agcctcccta	gtagctggga	ttatatgtgc	ccgcccacata	taattagccc	11100
cctggctgat	tttttttttt	tttttgtatt	tttagtagag	acagggtttc	gccatgttgg	11160
ccaggctgat	ctcgaaactc	tgacctcagg	tgatccactc	gctcggcctc	cccaaagtgc	11220
taggattaca	ggtgtgagtc	actgtgcccg	gccaagtttt	gcatttttag	tagactcccg	11280
gtctttaact	ccggacctca	ggtgatctgc	ctgccttggc	ctcccaaagt	gctggggtta	11340
caggcataag	ccattgtgct	cagccttata	tgcctatttt	taagagtttg	tgggtcaaaa	11400
tgagaccaat	gggaccattt	ttaaggaggc	aatccaaggg	cgagttggat	ggaactgaat	11460
taattgaacc	gaagttgggt	ttagacaagg	aactacaaga	tccctgaggc	atccctgtgt	11520
agaattgaga	tccacgcctt	ccaggacaag	gcttatggag	tgttaaaatg	aaagtgcctt	11580
gccactctga	caggcaatag	ctcttttgtc	ttggccttgg	ggtaataccg	ggggatggcg	11640
cttggccaga	aactgtcagt	tgccaacgag	aactcaagct	ggttcactgg	cagtccgaaa	11700
acagaaaaga	gcccgtggca	gtccctcacc	cctaagggca	aggacagcca	ggtatccctt	11760
ctctagggct	tcaggatccc	acagaagagc	tgcctccacc	gggaaccggca	gttccccaaa	11820
gagtaaagaa	ccagaccgtg	gaaggaagca	gagagaaaaa	ggaagaggga	aatcccagtg	11880
aagtccccgt	atgggccacc	aagatgccag	gcgaggtgtc	agagctccgg	aaccgggaag	11940
tggttggctc	ccgggtggta	aaagaactta	tcaacaaccg	tgtaggtctg	aaaaggaaaag	12000
ttttattaga	cggaaaggac	gaggcagcag	agcgcagtag	gcgcttcagc	aagagaggac	12060
tgagctccct	gcggggaact	gcagggtaat	ttggaccaca	ttagtcactt	aggtcatggt	12120
aaatggttac	atttgtcgat	attttggtgc	cttgatgtca	gcaaagtttg	cacaatgggt	12180
cttaacgtgc	actcattccg	gaaacgtaca	gaaattctag	ttacttataa	attcttggga	12240
cggaagcttg	gtaccagatg	tggctttaga	caatagggaa	gtgtcattct	gaattgctca	12300
gataaggggc	tttgccctct	gttggctcgac	ttgatggcca	ccaggtgata	tctgggtctct	12360
tcagtgtggc	tttgcagact	ataaaggcgc	agcgcgccaa	cgaggcggtt	tggccccaga	12420
cggcgagag	gaagggcaga	gtcggcggtc	ctgagacttg	gggcggcccc	ttggaggtca	12480
gccccgctcg	ctcctcccg	ccctctcctc	ctctccgagg	tccgaggcgg	gcagcgggct	12540
gtgggcgggc	aggaggtctg	ggagggggcg	ggggcaggaa	ggggcggggg	gctcggcgca	12600
ctcggcagga	agagaccgac	ccgccaccgc	ccgtagcccg	cgcgccccctg	gcactcaatc	12660
cccgccatgt	gggggtctct	gctcgccttg	gcgcctctcg	cgcgcggcgt	cggccccgct	12720
ctggggggcg	ccaggaaactc	ggtgctgggc	ctcgcgcagc	ccgggaccac	caaggtccca	12780
ggctcgaccc	cggccctgca	tagcagcccg	gcacagccgc	cggcgagagc	agctaaccgt	12840
gagttccccg	accgacggtc	cgtctccccg	caagccgact	gcccggctct	cctgccccgt	12900
ggggcgatcc	ctccctaaca	cgcgggcaca	cgcacaccca	cgcacactca	cagtcattgca	12960
cactcacccc	gcacgcacac	tgcactcac	gcgcacacac	gcgcgcgcac	tcacacacat	13020
tcacacacgc	gcacacttgc	actcacacgc	gcgcgcattc	acacgcattg	acacacacgc	13080
acactcacac	gcgcgtgcgc	gcacacacag	tgacgcgcgc	cgcacactca	cactcacagt	13140
gcacacacac	atatacacac	tcacactccc	tcaactccct	gctgggagca	atggctgctg	13200
actcggcagc	cccagttccc	tgcacagcct	agtcagcagt	cccaggacag	gcgccagtg	13260
gatgctgcct	cttccaagcc	ccaaaccttc	ccttttcacc	aaagacaaaa	caggccagaa	13320

ctggcaggag	gggagacaga	ggggcagaag	ctctcaaggt	gcagagcaag	actgogtagg	13380
agagagtttg	aaggcgaggg	ctggagagaa	agaaacaaaag	gaaagaaggg	agagpocctc	13440
gctgaggctg	cggggaggat	ggggcagagc	gggaagaggaa	ggcagcccga	cctccagct	13500
ttccagatgt	ggaataggag	aggaggagcg	caagcggagg	gcactcaggg	gcttctagag	13560
gaggcaagtg	gaggagggtc	ttgaagggtg	atgtccccga	gtcaggggag	tctggagaga	13620
gagagagaga	gagggtgccc	aagaaggaag	cggcgggcaa	aggcacaggg	gcaaccagatg	13680
cggaaatggg	cagcctgttc	tggaggcagc	tgtggagctt	cgatgggtac	ccccagcacc	13740
tgcctgggca	gagccttgtg	ctgaaggggc	ggcgggcagg	cccagccctg	aaagcctcga	13800
cacccaggca	gacatggatt	ccaggacagg	ccatctgagc	ccagagagca	gacacacaaa	13860
tggaaaggcg	acaggggttt	tggggcatga	tgtctgagct	ggagctaaga	aagcctcctt	13920
ggaaaggcat	ctgggtctgag	atgcaaaggga	agaatgggaa	ttaggtgaaa	aaatcagagg	13980
cgaggggtag	cattacaggg	gaggggatag	ctagtgcaga	ggcccggagg	taaagtgcga	14040
gactcagctc	tttggagcaa	ccgaacagtt	tctagaggct	gggtgcagct	ctccatttga	14100
ttagaggttc	acaggggagg	ctggccaagc	atgtagttaac	atcagggagg	agaaggagga	14160
gccaaaggaag	tgactggaga	ggcaggtttg	ggtcagattg	caggcctttg	atgtcctgtg	14220
aaggctgtta	gatacctggtg	gtgtggcctg	ctgtgggctc	acatgtcttc	ttgggtggc	14280
agacctttcc	atccgggggt	tcaccattct	tcctttcccc	catgctgtgc	ctctcggacc	14340
ccaagggacc	tcagaacagc	atgtccggat	tcgagtcac	aagaagaaaa	aggtcattat	14400
gaagaagcgg	aagaagctaa	ctctaactcg	ccccacccca	ctggtgactg	ccgggcccct	14460
tgtgaccccc	actccagcag	ggacccctcga	ccccgctgag	aaacaagaaa	caggtacttc	14520
ctctccaggg	gcccagccca	gacttgcagc	ccctggggca	ctttaccagc	acagctcttg	14580
gcctcatggg	caccggcacc	ccccttgctt	gcctagcgca	ggagcaacct	taggtctcagc	14640
ttcccacctg	ccctggctac	cctccctctg	gtcctgtctc	actgttctat	ccccgcccc	14700
ggctgtcctc	ctttgggtct	ggagtccctg	caggtttcag	atagccggct	tgaggcatcc	14760
agcagccagt	cctttgggtct	tggaccacac	cagggacggc	tcaacattca	ggtcagtaat	14820
cctggctcgg	agccatggtc	tcagggtagg	gaaggcagcc	cctgggagct	tctctcctgc	14880
ctcctctctg	tcctggcctg	ccccactctg	tccaactggg	cctgaccacc	atgtcctgtg	14940
tctgcagtea	ggcctggagg	acggcgatct	atatgatgga	gcctgggtgtg	ctgaggagca	15000
ggacgcagat	ccatggtttc	aggtggacgc	tgggcacccc	acccgcttct	cgggtgttat	15060
cacacagggc	aggaactctg	tctggagggtg	aggcagacta	accctaggtc	aggaggtcac	15120
agaaggactg	gggtgggagt	cctgggggca	ccgatgatct	ctctccacct	ctcctgccag	15180
gtatgactgg	gtcacatcat	acaaggctca	gttcagcaat	gacagtcgga	cctggtgggg	15240
aagtaggaac	cacagcagtg	ggatggacgc	agtgagtgg	cccactgtgg	ctggggccctc	15300
catgctggga	gttgggcacc	cagtcacaggc	taggtctgagg	ctcctctgag	gacaaggaa	15360
agacgccagc	ttaggtctcc	cagggggggtg	tggcttggtg	tcaagagggt	ggcacacggc	15420
aggcaccatt	gggagccagc	tgttttggga	catgccca	tcctccccag	ataatgccac	15480
caacgggtgg	gtgtctgttc	acgggtacagc	ttcctcctgg	cgtgcccctt	ctggcccggg	15540
gcctctggtc	cacatcactt	cttgccctct	cgtggttctg	acttcgcgat	ctcatggacc	15600
tctttttaca	gcaggctaca	atgtggagtc	ctggccagct	ctaggattgg	cttcccccca	15660
gtcatgtggc	caaaactggtc	taatgaactg	tgtccaatcc	agagagcaag	gctgcctagg	15720
gctgccatt	ggcaggggct	gtgggcggg	gtctgtgttt	gatgcacagt	gcaagtctct	15780
agctgagccc	actaggggtg	ggagacagta	agcttgagg	cctgagctcc	ttccctgggt	15840
cctgggccag	gcttctgggg	tttgagcagc	cacaacagag	aacttgcctg	ccccaggtat	15900
ttcctgccaa	ttcagaccca	gaaaactccag	tgtgaaacct	cctgccggag	ccccaggtgg	15960
ccgcttcat	tgcctgtctg	ccccagacct	ggctccaggg	aggegcgcct	tgcctccggg	16020
cagagatcct	ggcctgccca	gtctcaggtg	ggcagtcagg	ccagggttgg	ttgggcaggg	16080
cttggtatgca	gggtgcaccc	ttcactgtgg	acacaccctt	taccataaac	tcaacctcca	16140
ccagacccca	atgacctatt	ccttgaggcc	cctgcgtcgg	gatacctctga	ccctctagac	16200

ttccagcacc acaattacaa ggccatgagg aaggtcagat ataaccoccta tgacctggga 16260  
 aggaggggcc acccatctca ggcccccttc ccacccctcc accgggggac aacctgctgt 16320  
 gactgcgctt gtatgcccct gctgcccctt gatgtctcag ccttctctcc tgtggacccc 16380  
 taagctccat cccactttcc cttattatgg cgcaccccca gtccctaccc ttccctcccg 16440  
 ctctgctgcc gctcccccct tgtaccatga tgggatgccc cctctgtgtg ggccatcgtt 16500  
 gaactttttaa gtctttccat ggccatgtg atctgcccct ggggtgtacc ctcccatgcc 16560  
 tcatgccacg ctacactctg cccaccagct gatgaagcag gtacaagagc aatgccccaa 16620  
 catcaccgcc atctacagca ttgggaagag ctaccagggc ctgaagctgt atgtgatgga 16680  
 aatgtcggac aagcctgggg agcatgagct gggtaactgg atggggagtg gggagaggta 16740  
 ggccacagggc agggccccag gcctgaaccc gctgcaagcc cccatgtgtc cccaggggag 16800  
 cctgaggtgc gctacgtggc tggcatgcct gggaaagcag cctgggggag ggagttgctt 16860  
 ctgctcctga tgcagttcct gtgcccagag ttccctgagag ggaacccacg ggtgaccgag 16920  
 ctgctctctg agatgcgcat tccctgctg ccctccatga accctgatgg ctatgagatc 16980  
 gctacacacc gggtaggcca cccagcatga gggccactct gtccttctgc cctggtggct 17040  
 ggacctgctc gaattgaaca agcctcttgc ccggcagggt tcagagctgg tgggctgggc 17100  
 cgaggggcgc tggaaacaac agagcatcga tcttaacccat aattttgctg acctcaacac 17160  
 accactgtgg gaagcacagg acgatgggaa ggtgccccac atcgtcccca accatcacct 17220  
 gccattgccc acttaactaca cctgccccaa tgccaccgtg agtattttga gggcggcagt 17280  
 ggaggtctgt gggggggcga ccttgtctct gtctcctgcc cctcctgacc tgccccatcc 17340  
 aggtggtctc tgaacgcggg gcagtaatac agtggaatga ggggatcccc tttgtgctaa 17400  
 gtgccaacct ccacgggggt gagctcgtgg tgtccacccc attcgacatg actcgacccc 17460  
 cgtgggctgc ccgcgagctc acccccacac cagatgatgc tgtgtttcgc tggctcagca 17520  
 ctgtctatgc tggcagtaat ctggccatgc aggcacaccag ccgcgcgacc tgccacagcc 17580  
 aggacttctc cgtgcacggc aacatcatca accggggtga ctggcacacg gtccccggga 17640  
 gtatgtgctt gaggggtggg ttagccctgg ccccgtaacc cccgccctga taagacagcc 17700  
 tgcggttgcg tacagtgtct gctctgttcc cactctgaa gtgtccctca gagaaggag 17760  
 ggtagcggga ggatgggacc gcctcccgcc tgccttaggca gcagtgtctg tggccccctt 17820  
 aggcatgaat gacttcagct acctacacac caactgcttt gaggtcactg tggagctgtc 17880  
 ctgtgacaag ttccctcacc agaataaatt gccccaggag tgggagaaca acaaagacgc 17940  
 cctcctcacc taactggagc aggtcggatc tgcgtcccgg ccccccagct gcctgaatca 18000  
 ctctgctgtt ccatttaggc tacagctcct accagggggt cttctaaggt ccagctgagc 18060  
 attcagactc acaagatgcc atgggcccag cttgggtatca gattgtcttg gaagcacaca 18120  
 ggacaggaag tgcagtttgc tggcagcgtg gcctcgtgtt agagccgggt ggaggagcct 18180  
 ccattgcagt ctagggtgtg gtccgtggcg ctgccccaga gctatcctca ggagagactc 18240  
 acgtgaggca ggtgcaggag ctgtcctggc atagaagctt catgttccat ggagctcata 18300  
 acccttgtaa tagctccata agcagagctt ccaaagggtc taccaaagac aagcccaata 18360  
 acctgggaaa gcccaggat agataagcct tccctaccag tattttatcat tttcttagtc 18420  
 cagatgtgat ttgtcaatca ggatttcttt ttttttttcc ttccagaagt agtgtcacct 18480  
 aggaacacag tagacctacc actttgctca ggtttgcagg gcaacagagc cagcaagtta 18540  
 gctaaacagc acattatcct gccgaagggt aagggtctctg ataacctctt cccacacagg 18600  
 tgcgcattgg cattgcagga gtggtgaggg acaaggacac ggagcttggg attgctgacg 18660  
 ctgtcattgc cgtggatggg attaaccatg acgtgaccac ggggtgtgtt gaccgggagg 18720  
 gcaagggaag gggctggagg gctggaggct cgggaagaag cagaagatca ttaattgggt 18780  
 cctgatcgtg ccttccactc tccctcagct ggggcgggga ttattggcgt ctgctgaccc 18840  
 caggggacta catggtgact gccagtgcgc agggctacca ttccagtaca cggaaactgtc 18900  
 gggtcacctt tgaagagggc ccttccccct gcaatttctg gctcaccaag actcccaaac 18960  
 agaggctgag cagctgctg gcagctgggg ccaagggtgc cccggacctt cgcaggcgcc 19020  
 tggagcggct aaggggacag aaggattgat acctgcggtt taagagccct agggcaggct 19080

ggacctgtca	agacgggaag	gggaagagta	gagagggagg	gacaaagtga	ggaaaaggtg	19140
ctcattaaag	ctaccgggca	ccttagctca	tcttcgtgtt	gtctctgtgc	cccaggctct	19200
ccccccgggg	gcgggcctcg	gcccagccct	cagttcctat	tctgcacact	tgcacactct	19260
catcagtttg	cttctggaca	catttgtgtga	aaagaggatc	ccacctgggc	tcttcttgaa	19320
ccaagggcct	ggcagagcaa	ctcatttctt	ctgatcagct	tctgctacag	gtaccattac	19380
actgctgcca	ggcattctgt	aagcgctgc	tcattgccag	gtgtgcaagg	aatcaggatc	19440
agccgtgcct	gcactcaaac	tcctggggct	cctagtcaag	ggaaaggaca	gttcggtaca	19500
ttgtgagaca	tgttaggggtg	gaggccaggt	gcctgagag	tgcaggggag	ctgcacacgt	19560
gaaatacagc	actgcacatc	aacaggactg	gggcagtcaa	ggatgcaata	gaagtagtgg	19620
ctctagaagt	tcaggcgggga	ggtgggcagg	gtgtggagta	tggacagggga	tggctccaag	19680
gaggaggggtc	agccaaaggt	gggtcagctg	agaacatttg	aatttgcttc	agccattctc	19740
agagtattga	taactgatag	gctttgctga	gtttctatca	gaactgaagg	gaagtttgtg	19800
atcagtctgt	gtcttgccag	gtaaacaacc	cattctaggc	acttaaagtg	gagggaaatt	19860
taatgctgga	aattggatag	gaaggtgttg	gaagagctgg	atgaggccgg	gtgtggtggc	19920
tcacacctgt	aatcccagca	ctttgggagg	ctgaggtggg	aggattgctt	gagcccagga	19980
gtttgagacc	agcctggata	acatagccaa	accccgctc	tacaaaaata	agaaataaga	20040
aacatagcca	gctgtagtgg	cgcattggcta	agggaggcag	aggcaggagg	atcactggag	20100
cctgggaggt	ggaggctgca	gaggcagcag	tgagccatga	tggcgccact	atactccaac	20160
ctggtatggtc	ataacaaaat	aaacaaaaaa				20190

<210> 4

<211> 2205

<212> DNA

<213> Homo sapiens

<400> 4

atgtggggggc	tctgctcgc	cctggccgcc	ttcgccggcg	cctgcccgc	ggctctgggg	60
gcgcccagga	actcggtgct	gggcctcgcg	cagcccggga	ccaccaaggt	cccaggctcg	120
accccgggccc	tgcatagcag	ccggcacag	ccgcccggcg	agacagctaa	cgggacctca	180
gaacagcatg	tcggattcg	tgtcatcaag	aagaaaaagg	tcattatgaa	gaagcggaag	240
aagctaactc	taactcgccc	cacccactg	gtgactgcg	ggcccttgt	gaccccaact	300
ccagcagggga	ccctcgaccc	cgtgagaaa	caagaaacag	gctgtcctcc	tttgggtctg	360
gagtccttgc	gagtttcaga	tagccggctt	gaggcatcca	gcagccagtc	ctttggtctt	420
ggaccacacc	gaggacggct	caacattcag	tcaggcctgg	aggacggcga	tctatatgat	480
ggagcctggt	gtgctgagga	gcaggacgcc	gatccatggt	ttcagggtgga	cgtgggcac	540
cccaccgcct	tctcggtgt	tatcacacag	ggcaggaact	ctgtctggag	gtatgactgg	600
gtcacatcat	acaaggtcca	gttcagcaat	gacagtggga	cctggtgggg	aagtaggaac	660
cacagcagtg	ggatggacgc	agtatttctt	gccaatcag	acccagaaac	tccagtgtg	720
aacctcctgc	cggagcccca	ggtggccgc	ttcatcgcc	tgtgccccca	gacctggctc	780
cagggaggcg	cgccttgctt	ccgggcagag	atcctggcct	gcccagtcct	agaccccaat	840
gacctattcc	ttgaggcccc	tgcgtcgga	tcctctgacc	ctctagactt	tcagcatcac	900
aattacaagg	ccatgaggaa	gctgatgaag	caggtacaag	agcaatgccc	caacatcacc	960
cgcactctaca	gcattgggaa	gagctaccag	ggcctgaagc	tgtatgtgat	ggaaatgtcg	1020
gacaagcctg	gggagcatga	gctgggggag	cctgaggtgc	gtacgtggc	tggcatgcat	1080
gggaacgagg	ccctggggcg	ggagttgctt	ctgctcctga	tgcagttcct	gtgccaatgag	1140
ttcctgcgag	ggaacccacg	ggtgacccgg	ctgctctctg	agatgcgcct	tcacctgctg	1200
ccctccatga	accctgatgg	ctatgagatc	gcctaccacc	gggggttcaga	gctgggtgggc	1260

tgggccgagg	gocgctggaa	caaccagagc	atcgatotta	accataatit	tgetgacctc	1320
aacacaccac	tgtgggaagc	acaggacgat	gggaagggtc	cccacatcgt	ccccaccat	1380
cacctgccat	tgccactta	ctacacccctg	cccaatgcc	ccgtggctcc	tgaaacgcgg	1440
gcagtaata	agtggatgaa	gcggatcccc	tttgtgttaa	gtgccaaact	ccacgggggt	1500
gagctcgtgg	tgtctaccc	attcgacatg	actcgacccc	cgtgggctgc	ccgcgagctc	1560
acgcccacac	cagatgatgc	tgtgtttcgc	tggtcagca	ctgtctatgc	tggcagtaat	1620
ctggccatgc	aggacaccag	ccgcgcagccc	tgccacagcc	aggactttct	cgtgcacggc	1680
aacatcatca	acggggctga	ctggcacacg	gtccccggga	gcattgaatga	cttcagctac	1740
ctacacacca	actgctttga	ggtcaactgtg	gagctgtcct	gtgacaagtt	ccctcacgag	1800
aatgaattgc	cccaggagtg	ggagaacaac	aaagacgccc	tcttcaccta	cctggagcag	1860
gtgcgcattg	gcattgcagg	agtgggtgag	gacaaggaca	cggagcttgg	gattgctgac	1920
gtgtgcattg	ccgtggatgg	gattaacct	gacgtgacca	cggcgtgggg	cggggattat	1980
tggcgtctgc	tgaccccagg	ggactacatg	gtgactgcca	gtgcgcaggg	ctaccattca	2040
gtgacacgga	actgtcgggt	cacctttgaa	gagggcacct	tccctgcaa	tttcgtgctc	2100
accaagactc	ccaaacagag	gctgcgcgag	ctgctggcag	ctggggccaa	ggtgcccccg	2160
gaccttcgca	ggcgcctgga	gcggctaagg	ggacagaagg	attga		2205

<210> 5  
 <211> 1725  
 <212> DNA  
 <213> Homo sapiens

atgtgggggc	tctgtctcgc	cctggccgcc	ttcgcgcgcg	ccgtcggccc	ggctctgggg	60
ggcccaggga	actcgggtgt	gggcctcgcg	cagcccggga	ccaccaaggt	cccaggctcg	120
accccggccc	tgcatagcag	cccgccacag	ccgcgcggcg	agacagctaa	cgggacctca	180
gaacagcatg	tcgggattcg	agtcattcaag	aagaaaaagg	tcattatgaa	gaagcgggaag	240
aagctaactc	taactcgcgc	caccccactg	gtgactgcgc	ggccccttgt	gacccccact	300
ccagcaggga	ccctcgaccc	cgtcgagaaa	caagaaacag	gctgtcctcc	tttgggtctg	360
gagtcctctg	gagtttcaga	tagccggctt	gaggcatcca	gcagccagtc	ctttgggtctt	420
ggaccacacc	gaggacggct	caacattcag	tcaggccctgg	aggacggcga	tctatatgat	480
ggagcctggg	gtgctgagga	gcaggacgcc	gatccatggg	ttcagggtgga	cgtcgggcac	540
cccaccgcgt	tctcgggtgt	tatcacacag	ggcagggaact	ctgtctggag	gtatgactgg	600
gtcacatcat	acaagggtcca	gttcagcaat	gacagtcgga	cctgggtgggg	aagtaggaac	660
cacagcagtg	ggatggacgc	agtatttctt	gccaatcag	acccagaaac	tccagtgtctg	720
aacctcctgc	cggagcccca	ggtggccccc	ttcatctgcc	tgtgccccca	gacctggctc	780
caggggaggcg	cgccttgccct	ccgggcagag	atcctggcct	gcccagttct	agaccccaat	840
gacctattcc	ttgaggcccc	tgcgtcggga	tcctctgacc	ctctagaatt	tcagcatcac	900
aattacaagg	ccatgaggaa	gctgatgaag	caggtaacaag	agcaatgccc	caacatcacc	960
cgcattctaca	gcattgggaa	gagctaccag	ggcctgaagc	tgtatgtgat	ggaaatgtcg	1020
gacaagcctg	gggagcatga	gctgggggag	cctgagggtgc	gtaactgtgc	tggcatgcac	1080
gggaacgagg	ccctggggcg	ggagttgctt	ctgctcctga	tgcagttcct	gtgccatgag	1140
ttcctgcgag	ggaacccacg	ggtgaacccg	ctgctctctg	agatgcgcac	tcacctgctg	1200
ccctccatga	acctgatgg	ctatgagatc	gcctaccacc	gggggttcaga	gctggtgggc	1260
tgggccgagg	gocgctggaa	caaccagagc	atcgatotta	accataatit	tgetgacctc	1320
aacacaccac	tgtgggaagc	acaggacgat	gggaagggtc	cccacatcgt	ccccaccat	1380
cacctgccat	tgccactta	ctacacccctg	cccaatgcc	ccgtggctcc	tgaaacgcgg	1440

gcagtaatca agtggatgaa gccggtcccc ttgtgtctaa gtgccaaact ccacgggggt 1500  
gagctcgtgg tgtcctaccc attcgacatg gtgaactgcca gtgccgaggg ctaccattca 1560  
gtgacacgga actgtcgggt cacttttgaa gagggcccct tccctgcaa ttctgtgtc 1620  
accaagactc ccaaacagag gctgcgcgag ctgtggcag ctggggccaa ggtgcccccg 1680  
gaccttcgca gccgcctgga gccgctaagg ggacagaagg attga 1725

<210> 6  
<211> 574  
<212> PRT  
<213> Homo sapiens

<400> 6  
Met Trp Gly Leu Leu Leu Ala Leu Ala Ala Phe Ala Pro Ala Val Gly  
1 5 10 15  
Pro Ala Leu Gly Ala Pro Arg Asn Ser Val Leu Gly Leu Ala Gln Pro  
20 25 30  
Gly Thr Thr Lys Val Pro Gly Ser Thr Pro Ala Leu His Ser Ser Pro  
35 40 45  
Ala Gln Pro Pro Ala Glu Thr Ala Asn Gly Thr Ser Glu Gln His Val  
50 55 60  
Arg Ile Arg Val Ile Lys Lys Lys Lys Val Ile Met Lys Lys Arg Lys  
65 70 75 80  
Lys Leu Thr Leu Thr Arg Pro Thr Pro Leu Val Thr Ala Gly Pro Leu  
85 90 95  
Val Thr Pro Thr Pro Ala Gly Thr Leu Asp Pro Ala Glu Lys Gln Glu  
100 105 110  
Thr Gly Cys Pro Pro Leu Gly Leu Glu Ser Leu Arg Val Ser Asp Ser  
115 120 125  
Arg Leu Glu Ala Ser Ser Ser Gln Ser Phe Gly Leu Gly Pro His Arg  
130 135 140  
Gly Arg Leu Asn Ile Gln Ser Gly Leu Glu Asp Gly Asp Leu Tyr Asp  
145 150 155 160  
Gly Ala Trp Cys Ala Glu Glu Gln Asp Ala Asp Pro Trp Phe Gln Val  
165 170 175  
Asp Ala Gly His Pro Thr Arg Phe Ser Gly Val Ile Thr Gln Gly Arg  
180 185 190



Asn Ser Val Trp Arg Tyr Asp Trp Val Thr Ser Tyr Lys Val Gln Phe  
 195 200 205

Ser Asn Asp Ser Arg Thr Trp Trp Gly Ser Arg Asn His Ser Ser Gly  
 210 215 220

Met Asp Ala Val Phe Pro Ala Asn Ser Asp Pro Glu Thr Pro Val Leu  
 225 230 235 240

Asn Leu Leu Pro Glu Pro Gln Val Ala Arg Phe Ile Arg Leu Leu Pro  
 245 250 255

Gln Thr Trp Leu Gln Gly Gly Ala Pro Cys Leu Arg Ala Glu Ile Leu  
 260 265 270

Ala Cys Pro Val Ser Asp Pro Asn Asp Leu Phe Leu Glu Ala Pro Ala  
 275 280 285

Ser Gly Ser Ser Asp Pro Leu Asp Phe Gln His His Asn Tyr Lys Ala  
 290 295 300

Met Arg Lys Leu Met Lys Gln Val Gln Glu Gln Cys Pro Asn Ile Thr  
 305 310 315 320

Arg Ile Tyr Ser Ile Gly Lys Ser Tyr Gln Gly Leu Lys Leu Tyr Val  
 325 330 335

Met Glu Met Ser Asp Lys Pro Gly Glu His Glu Leu Gly Glu Pro Glu  
 340 345 350

Val Arg Tyr Val Ala Gly Met His Gly Asn Glu Ala Leu Gly Arg Glu  
 355 360 365

Leu Leu Leu Leu Leu Met Gln Phe Leu Cys His Glu Phe Leu Arg Gly  
 370 375 380

Asn Pro Arg Val Thr Arg Leu Leu Ser Glu Met Arg Ile His Leu Leu  
 385 390 395 400

Pro Ser Met Asn Pro Asp Gly Tyr Glu Ile Ala Tyr His Arg Gly Ser  
 405 410 415

Glu Leu Val Gly Trp Ala Glu Gly Arg Trp Asn Asn Gln Ser Ile Asp  
 420 425 430

Leu Asn His Asn Phe Ala Asp Leu Asn Thr Pro Leu Trp Glu Ala Gln  
 435 440 445

Asp Asp Gly Lys Val Pro His Ile Val Pro Asn His His Leu Pro Leu  
 450 455 460

Pro Thr Tyr Tyr Thr Leu Pro Asn Ala Thr Val Ala Pro Glu Thr Arg  
 465 470 475 480

Ala Val Ile Lys Trp Met Lys Arg Ile Pro Phe Val Leu Ser Ala Asn  
 485 490 495

Leu His Gly Gly Glu Leu Val Val Ser Tyr Pro Phe Asp Met Val Thr  
 500 505 510

Ala Ser Ala Glu Gly Tyr His Ser Val Thr Arg Asn Cys Arg Val Thr  
 515 520 525

Phe Glu Glu Gly Pro Phe Pro Cys Asn Phe Val Leu Thr Lys Thr Pro  
 530 535 540

Lys Gln Arg Leu Arg Glu Leu Leu Ala Ala Gly Ala Lys Val Pro Pro  
 545 550 555 560

Asp Leu Arg Arg Arg Leu Glu Arg Leu Arg Gly Gln Lys Asp  
 565 570

<210> 7  
 <211> 1972  
 <212> DNA  
 <213> Homo sapiens

<400> 7  
 atgtgggggc tectgctcgc cctggccgcc ttgcgcgcgg ccgtcggccc ggctctgggg 60  
 gcgcccagga actcgggtgct gggcctcgcg cagcccggga ccaccaaggt cccaggctcg 120  
 accccggccc tgcatagcag cccggcacag ccgcccggcg agacagctaa cgggacctca 180  
 gaacagcatg tccggattcg agtcatcaag aagaaaaagg tcattatgaa gaagcgggaag 240  
 aagctaactc taactcgccc caccocactg gtgactgccg ggcccccttgt gacccccact 300  
 ccagcaggga ccctcgaccc cgctgagaaa caagaaacag gctgtcctcc tttgggtctg 360  
 gagtccctgc gagtttcaga tagccggctt gaggcattcca gcagccagtc ctttgggtctt 420  
 ggaccacacc gaggaaggct caacattcag tcaggcctgg aggacggcga tctatatgat 480  
 ggagcctggt gtgctgagga gcaggacgcc gatccatggt ttcagggtgga cgctgggcac 540  
 cccaccgcgt tctcgggtgt tatcacacag ggcagagatc ctggcctgcc cagtctcaga 600  
 ccccaatgac ctattccttg aggccctgc gtcgggatcc tctgaccctc tagactttca 660  
 gcatcacaat tacaaggcca tgaggaagct gatgaagcag gtacaagagc aatgccccaa 720  
 catcaccgcg atctacagca ttgggaagag ctaccagggc ctgaagctgt atgtgatgga 780  
 aatgtcggac aagcctgggg agcatgagct gggggagcct gaggtgcgct acgtggctgg 840  
 catgcatggg aacgaggccc tggggcgga gttgcttctg ctctgatgc agttcctgtg 900  
 ccatgagttc ctgcgaggga acccacgggt gaccggctg ctctctgaga tgcgcattca 960

```

cctgctgccc tccatgaacc ctgatggcta tgagatcgcc taaccacggg gttcagagct 1020
ggtgggctgg gcgaggggccc gctggaacaa ccagagcacc gatcttaacc ataattttgc 1080
tgacctcaac acaccactgt gggaagcaca ggaagatggg aagggtgccc acatcgcccc 1140
caaccatcac ctgccattgc ccacttacta caccctgccc aatgcccacg tggctcctga 1200
aacgggggca gtaatcaagt ggatgaagcg gatccccctt gtgctaagtg ccaacctcca 1260
cgggggtgag ctgctgggtg cctacccatt cgacatgaat cgcaccccggt gggctgcccc 1320
cgagctcacg cccacaccag atgatgctgt gtttcgctgg ctacgcaactg tctatgctgg 1380
cagtaatctg gccatgcagg acaccagcgg ccgacccctg ccacgcccagg acttctccgt 1440
gcacggcaac atcatcaacg gggctgactg gcacacggtc cccgggagca tgaatgactt 1500
cagctaccta cacaccaact gctttgaggt cactgtggag ctgtcctgtg acaagttccc 1560
tcacgagaat gaattgcccc aggagtggga gaacaacaaa gacgcctccc tcacctacct 1620
ggagcaggtg cgcctgggca ttgcaggagt ggtgagggac aaggacaagg agcttgggat 1680
tgctgaagct gtcattgccg tggatgggat taaccatgac gtgaccaagg cgtggggcgg 1740
ggattattgg cgtctgctga cccaggggga ctacatggtg actgccagtg ccgagggcta 1800
ccattcagtg acacggaact gtcgggtcac ctttgaagag ggccccctcc cctgcaattt 1860
cgtgctcacc aagactccca aacagaggct gcgcgagctg ctggcagctg gggccaaggt 1920
gcccccgga ccttcgaggc gcctggagcg gctaagggga ccagaaggatt ga 1972

```

<210> 8

<211> 202

<212> PRT

<213> Homo sapiens

<400> 8

```

Met Trp Gly Leu Leu Leu Ala Leu Ala Ala Phe Ala Pro Ala Val Gly
  1              5              10             15

```

```

Pro Ala Leu Gly Ala Pro Arg Asn Ser Val Leu Gly Leu Ala Gln Pro
      20              25             30

```

```

Gly Thr Thr Lys Val Pro Gly Ser Thr Pro Ala Leu His Ser Ser Pro
      35              40             45

```

```

Ala Gln Pro Pro Ala Glu Thr Ala Asn Gly Thr Ser Glu Gln His Val
      50              55             60

```

```

Arg Ile Arg Val Ile Lys Lys Lys Lys Val Ile Met Lys Lys Arg Lys
      65              70             75             80

```

```

Lys Leu Thr Leu Thr Arg Pro Thr Pro Leu Val Thr Ala Gly Pro Leu
      85              90             95

```

```

Val Thr Pro Thr Pro Ala Gly Thr Leu Asp Pro Ala Glu Lys Gln Glu
      100             105            110

```

```

Thr Gly Cys Pro Pro Leu Gly Leu Glu Ser Leu Arg Val Ser Asp Ser
      115             120            125

```

Arg Leu Glu Ala Ser Ser Ser Gln Ser Phe Gly Leu Gly Pro His Arg  
130 135 140

Gly Arg Leu Asn Ile Gln Ser Gly Leu Glu Asp Gly Asp Leu Tyr Asp  
145 150 155 160

Gly Ala Trp Cys Ala Glu Glu Gln Asp Ala Asp Pro Trp Phe Gln Val  
165 170 175

Asp Ala Gly His Pro Thr Arg Phe Ser Gly Val Ile Thr Gln Gly Arg  
180 185 190

Asp Pro Gly Leu Pro Ser Leu Arg Pro Gln  
195 200

<210> 9

<211> 719

<212> PRT

<213> Mus musculus

<400> 9

Met Gln Ala Gly Ala Asn Glu Asp Asp Tyr Tyr Asp Gly Ala Trp Cys  
1 5 10 15

Ala Glu Asp Glu Ser Gln Thr Gln Trp Ile Glu Val Asp Thr Arg Arg  
20 25 30

Thr Thr Arg Phe Thr Gly Val Ile Thr Gln Gly Arg Asp Ser Ser Ile  
35 40 45

His Asp Asp Phe Val Thr Thr Phe Phe Val Gly Phe Ser Asn Asp Ser  
50 55 60

Gln Thr Trp Val Met Tyr Thr Asn Gly Tyr Glu Glu Met Thr Phe Tyr  
65 70 75 80

Gly Asn Val Asp Lys Asp Thr Pro Val Leu Ser Glu Leu Pro Glu Pro  
85 90 95

Val Val Ala Arg Phe Ile Arg Ile Tyr Pro Leu Thr Trp Asn Gly Ser  
100 105 110

Leu Cys Met Arg Leu Glu Val Leu Gly Cys Pro Val Thr Pro Val Tyr  
115 120 125

Ser Tyr Tyr Ala Gln Asn Glu Val Val Thr Thr Asp Ser Leu Asp Phe

130	135	140
Arg His His Ser Tyr Lys Asp Met Arg Gln Leu Met Lys Ala Val Asn		
145	150	155 160
Glu Glu Cys Pro Thr Ile Thr Arg Thr Tyr Ser Leu Gly Lys Ser Ser		
	165	170 175
Arg Gly Leu Lys Ile Tyr Ala Met Glu Ile Ser Asp Asn Pro Gly Asp		
	180	185 190
His Glu Leu Gly Glu Pro Glu Phe Arg Tyr Thr Ala Gly Ile His Gly		
	195	200 205
Asn Glu Val Leu Gly Arg Glu Leu Leu Leu Leu Met Gln Tyr Leu		
	210	215 220
Cys Gln Glu Tyr Arg Asp Gly Asn Pro Arg Val Arg Asn Leu Val Gln		
225	230	235 240
Asp Thr Arg Ile His Leu Val Pro Ser Leu Asn Pro Asp Gly Tyr Glu		
	245	250 255
Val Ala Ala Gln Met Gly Ser Glu Phe Gly Asn Trp Ala Leu Gly Leu		
	260	265 270
Trp Thr Glu Glu Gly Phe Asp Ile Phe Glu Asp Phe Pro Asp Leu Asn		
	275	280 285
Ser Val Leu Trp Ala Ala Glu Glu Lys Lys Trp Val Pro Tyr Arg Val		
	290	295 300
Pro Asn Asn Asn Leu Pro Ile Pro Glu Arg Tyr Leu Ser Pro Asp Ala		
305	310	315 320
Thr Val Ser Thr Glu Val Arg Ala Ile Ile Ser Trp Met Glu Lys Asn		
	325	330 335
Pro Phe Val Leu Gly Ala Asn Leu Asn Gly Gly Glu Arg Leu Val Ser		
	340	345 350
Tyr Pro Tyr Asp Met Ala Arg Thr Pro Ser Gln Glu Gln Leu Leu Ala		
	355	360 365
Glu Ala Leu Ala Ala Ala Arg Gly Glu Asp Asp Asp Gly Val Ser Glu		
	370	375 380
Ala Gln Glu Thr Pro Asp His Ala Ile Phe Arg Trp Leu Ala Ile Ser		

385		390		395		400
Phe Ala Ser Ala His Leu Thr Met Thr Glu Pro Tyr Arg Gly Gly Cys						
	405		410		415	
Gln Ala Gln Asp Tyr Thr Ser Gly Met Gly Ile Val Asn Gly Ala Lys						
	420		425		430	
Trp Asn Pro Arg Ser Gly Thr Phe Asn Asp Phe Ser Tyr Leu His Thr						
	435		440		445	
Asn Cys Leu Glu Leu Ser Val Tyr Leu Gly Cys Asp Lys Phe Pro His						
	450		455		460	
Glu Ser Glu Leu Pro Arg Glu Trp Glu Asn Asn Lys Glu Ala Leu Leu						
465		470		475		480
Thr Phe Met Glu Gln Val His Arg Gly Ile Lys Gly Val Val Thr Asp						
	485		490		495	
Glu Gln Gly Ile Pro Ile Ala Asn Ala Thr Ile Ser Val Ser Gly Ile						
	500		505		510	
Asn His Gly Val Lys Thr Ala Ser Gly Gly Asp Tyr Trp Arg Ile Leu						
	515		520		525	
Asn Pro Gly Glu Tyr Arg Val Thr Ala His Ala Glu Gly Tyr Thr Ser						
	530		535		540	
Ser Ala Lys Ile Cys Asn Val Asp Tyr Asp Ile Gly Ala Thr Gln Cys						
545		550		555		560
Asn Phe Ile Leu Ala Arg Ser Asn Trp Lys Arg Ile Arg Glu Ile Leu						
	565		570		575	
Ala Met Asn Gly Asn Arg Pro Ile Leu Arg Val Asp Pro Ser Arg Pro						
	580		585		590	
Met Thr Pro Gln Gln Arg Arg Met Gln Gln Arg Arg Leu Gln Tyr Arg						
	595		600		605	
Leu Arg Met Arg Glu Gln Met Arg Leu Arg Arg Leu Asn Ser Thr Ala						
	610		615		620	
Gly Pro Ala Thr Ser Pro Thr Pro Ala Leu Met Pro Pro Pro Ser Pro						
625		630		635		640
Thr Pro Ala Ile Thr Leu Arg Pro Trp Glu Val Leu Pro Thr Thr Thr						

	645		650		655
Ala Gly Trp Glu Glu Ser Glu Thr Glu Thr Tyr Thr Glu Val Val Thr					
	660		665		670
Glu Phe Glu Thr Glu Tyr Gly Thr Asp Leu Glu Val Glu Glu Ile Glu					
	675		680		685
Glu Glu Glu Glu Glu Glu Glu Glu Glu Met Asp Thr Gly Leu Thr Phe					
	690		695		700
Pro Leu Thr Thr Val Glu Thr Tyr Thr Val Asn Phe Gly Asp Phe					
	705		710		715
<210> 10					
<211> 1128					
<212> PRT					
<213> Mus musculus					
<400> 10					
Met Ala Pro Val Arg Thr Ala Ser Leu Leu Cys Gly Leu Leu Ala Leu					
1		5		10	15
Leu Thr Leu Cys Pro Glu Gly Asn Pro Gln Thr Val Leu Thr Asp Asp					
	20		25		30
Glu Ile Glu Glu Phe Leu Glu Gly Phe Leu Ser Glu Leu Glu Thr Gln					
	35		40		45
Ser Pro Pro Arg Glu Asp Asp Val Glu Val Gln Pro Leu Pro Glu Pro					
	50		55		60
Thr Gln Arg Pro Arg Lys Ser Lys Ala Gly Gly Lys Gln Arg Ala Asp					
	65		70		75
Val Glu Val Pro Pro Glu Lys Asn Lys Asp Lys Glu Lys Lys Gly Lys					
		85		90	95
Lys Asp Lys Gly Pro Lys Ala Thr Lys Pro Leu Glu Gly Ser Thr Arg					
	100		105		110
Pro Thr Lys Lys Pro Lys Glu Lys Pro Pro Lys Ala Thr Lys Lys Pro					
	115		120		125
Lys Glu Lys Pro Pro Lys Ala Thr Lys Lys Pro Lys Glu Lys Pro Pro					
	130		135		140

Lys Ala Thr Lys Lys Pro Lys Glu Lys Pro Pro Lys Ala Thr Lys Arg  
 145 150 155 160  
 Pro Ser Ala Gly Lys Lys Phe Ser Thr Val Ala Pro Leu Glu Thr Leu  
 165 170 175  
 Asp Arg Leu Leu Pro Ser Pro Ser Asn Pro Ser Ala Gln Glu Leu Pro  
 180 185 190  
 Gln Lys Arg Asp Thr Pro Phe Pro Asn Ala Trp Gln Gly Gln Gly Glu  
 195 200 205  
 Glu Thr Gln Val Glu Ala Lys Gln Pro Arg Pro Glu Pro Glu Glu Glu  
 210 215 220  
 Thr Glu Met Pro Thr Leu Asp Tyr Asn Asp Gln Ile Glu Lys Glu Asp  
 225 230 235 240  
 Tyr Glu Asp Phe Glu Tyr Ile Arg Arg Gln Lys Gln Pro Arg Pro Thr  
 245 250 255  
 Pro Ser Arg Arg Arg Leu Trp Pro Glu Arg Pro Glu Glu Lys Thr Glu  
 260 265 270  
 Glu Pro Glu Glu Arg Lys Glu Val Glu Pro Pro Leu Lys Pro Leu Leu  
 275 280 285  
 Pro Pro Asp Tyr Gly Asp Ser Tyr Val Ile Pro Asn Tyr Asp Asp Leu  
 290 295 300  
 Asp Tyr Tyr Phe Pro His Pro Pro Pro Gln Lys Pro Asp Val Gly Gln  
 305 310 315 320  
 Glu Val Asp Glu Glu Lys Glu Glu Met Lys Lys Pro Lys Lys Glu Gly  
 325 330 335  
 Ser Ser Pro Lys Glu Asp Thr Glu Asp Lys Trp Thr Val Glu Lys Asn  
 340 345 350  
 Lys Asp His Lys Gly Pro Arg Lys Gly Glu Glu Leu Glu Glu Glu Trp  
 355 360 365  
 Ala Pro Val Glu Lys Ile Lys Cys Pro Pro Ile Gly Met Glu Ser His  
 370 375 380  
 Arg Ile Glu Asp Asn Gln Ile Arg Ala Ser Ser Met Leu Arg His Gly  
 385 390 395 400



Leu Gly Ala Gln Arg Gly Arg Leu Asn Met Gln Ala Gly Ala Asn Glu  
 405 410 415  
 Asp Asp Tyr Tyr Asp Gly Ala Trp Cys Ala Glu Asp Glu Ser Gln Thr  
 420 425 430  
 Gln Trp Ile Glu Val Asp Thr Arg Arg Thr Thr Arg Phe Thr Gly Val  
 435 440 445  
 Ile Thr Gln Gly Arg Asp Ser Ser Ile His Asp Asp Phe Val Thr Thr  
 450 455 460  
 Phe Phe Val Gly Phe Ser Asn Asp Ser Gln Thr Trp Val Met Tyr Thr  
 465 470 475 480  
 Asn Gly Tyr Glu Glu Met Thr Phe Tyr Gly Asn Val Asp Lys Asp Thr  
 485 490 495  
 Pro Val Leu Ser Glu Leu Pro Glu Pro Val Val Ala Arg Phe Ile Arg  
 500 505 510  
 Ile Tyr Pro Leu Thr Trp Asn Gly Ser Leu Cys Met Arg Leu Glu Val  
 515 520 525  
 Leu Gly Cys Pro Val Thr Pro Val Tyr Ser Tyr Tyr Ala Gln Asn Glu  
 530 535 540  
 Val Val Thr Thr Asp Ser Leu Asp Phe Arg His His Ser Tyr Lys Asp  
 545 550 555 560  
 Met Arg Gln Leu Met Lys Ala Val Asn Glu Glu Cys Pro Thr Ile Thr  
 565 570 575  
 Arg Thr Tyr Ser Leu Gly Lys Ser Ser Arg Gly Leu Lys Ile Tyr Ala  
 580 585 590  
 Met Glu Ile Ser Asp Asn Pro Gly Asp His Glu Leu Gly Glu Pro Glu  
 595 600 605  
 Phe Arg Tyr Thr Ala Gly Ile His Gly Asn Glu Val Leu Gly Arg Glu  
 610 615 620  
 Leu Leu Leu Leu Leu Met Gln Tyr Leu Cys Gln Glu Tyr Arg Asp Gly  
 625 630 635 640  
 Asn Pro Arg Val Arg Asn Leu Val Gln Asp Thr Arg Ile His Leu Val  
 645 650 655

Pro	Ser	Leu	Asn	Pro	Asp	Gly	Tyr	Glu	Val	Ala	Ala	Gln	Met	Gly	Ser	660	665	670	
Glu	Phe	Gly	Asn	Trp	Ala	Leu	Gly	Leu	Trp	Thr	Glu	Glu	Gly	Phe	Asp	675	680	685	
Ile	Phe	Glu	Asp	Phe	Pro	Asp	Leu	Asn	Ser	Val	Leu	Trp	Ala	Ala	Glu	690	695	700	
Glu	Lys	Lys	Trp	Val	Pro	Tyr	Arg	Val	Pro	Asn	Asn	Asn	Leu	Pro	Ile	705	710	715	720
Pro	Glu	Arg	Tyr	Leu	Ser	Pro	Asp	Ala	Thr	Val	Ser	Thr	Glu	Val	Arg	725	730	735	
Ala	Ile	Ile	Ser	Trp	Met	Glu	Lys	Asn	Pro	Phe	Val	Leu	Gly	Ala	Asn	740	745	750	
Leu	Asn	Gly	Gly	Glu	Arg	Leu	Val	Ser	Tyr	Pro	Tyr	Asp	Met	Ala	Arg	755	760	765	
Thr	Pro	Ser	Gln	Glu	Gln	Leu	Leu	Ala	Glu	Ala	Leu	Ala	Ala	Ala	Arg	770	775	780	
Gly	Glu	Asp	Asp	Asp	Gly	Val	Ser	Glu	Ala	Gln	Glu	Thr	Pro	Asp	His	785	790	795	800
Ala	Ile	Phe	Arg	Trp	Leu	Ala	Ile	Ser	Phe	Ala	Ser	Ala	His	Leu	Thr	805	810	815	
Met	Thr	Glu	Pro	Tyr	Arg	Gly	Gly	Cys	Gln	Ala	Gln	Asp	Tyr	Thr	Ser	820	825	830	
Gly	Met	Gly	Ile	Val	Asn	Gly	Ala	Lys	Trp	Asn	Pro	Arg	Ser	Gly	Thr	835	840	845	
Phe	Asn	Asp	Phe	Ser	Tyr	Leu	His	Thr	Asn	Cys	Leu	Glu	Leu	Ser	Val	850	855	860	
Tyr	Leu	Gly	Cys	Asp	Lys	Phe	Pro	His	Glu	Ser	Glu	Leu	Pro	Arg	Glu	865	870	875	880
Trp	Glu	Asn	Asn	Lys	Glu	Ala	Leu	Leu	Thr	Phe	Met	Glu	Gln	Val	His	885	890	895	
Arg	Gly	Ile	Lys	Gly	Val	Val	Thr	Asp	Glu	Gln	Gly	Ile	Pro	Ile	Ala	900	905	910	

Asn Ala Thr Ile Ser Val Ser Gly Ile Asn His Gly Val Lys Thr Ala  
915 920 925

Ser Gly Gly Asp Tyr Trp Arg Ile Leu Asn Pro Gly Glu Tyr Arg Val  
930 935 940

Thr Ala His Ala Glu Gly Tyr Thr Ser Ser Ala Lys Ile Cys Asn Val  
945 950 955 960

Asp Tyr Asp Ile Gly Ala Thr Gln Cys Asn Phe Ile Leu Ala Arg Ser  
965 970 975

Asn Trp Lys Arg Ile Arg Glu Ile Leu Ala Met Asn Gly Asn Arg Pro  
980 985 990

Ile Leu Gly Val Asp Pro Ser Arg Pro Met Thr Pro Gln Gln Arg Arg  
995 1000 1005

Met Gln Gln Arg Arg Leu Gln Tyr Arg Leu Arg Met Arg Glu Gln Met  
1010 1015 1020

Arg Leu Arg Arg Leu Asn Ser Thr Ala Gly Pro Ala Thr Ser Pro Thr  
1025 1030 1035 1040

Pro Ala Leu Met Pro Pro Pro Ser Pro Thr Pro Ala Ile Thr Leu Arg  
1045 1050 1055

Pro Trp Glu Val Leu Pro Thr Thr Thr Ala Gly Trp Glu Glu Ser Glu  
1060 1065 1070

Thr Glu Thr Tyr Thr Glu Val Val Thr Glu Phe Glu Thr Glu Tyr Gly  
1075 1080 1085

Thr Asp Leu Glu Val Glu Glu Ile Glu Glu Glu Glu Glu Glu Glu  
1090 1095 1100

Glu Glu Met Asp Thr Gly Leu Thr Phe Pro Leu Thr Thr Val Glu Thr  
1105 1110 1115 1120

Tyr Thr Val Asn Phe Gly Asp Phe  
1125

<210> 11

<211> 845

<212> PRT

<213> Homo sapiens

<400> 11

Met Asp Tyr Tyr Phe Gly Pro Pro Pro Pro Gln Lys Pro Asp Ala Glu  
1 5 10 15

Arg Gln Thr Asp Glu Glu Lys Glu Glu Leu Lys Lys Pro Lys Lys Glu  
20 25 30

Asp Ser Ser Pro Lys Glu Glu Thr Asp Lys Trp Ala Val Glu Lys Gly  
35 40 45

Lys Asp His Lys Glu Pro Arg Lys Gly Glu Glu Leu Glu Glu Glu Trp  
50 55 60

Thr Pro Thr Glu Lys Val Lys Cys Pro Pro Ile Gly Met Glu Ser His  
65 70 75 80

Arg Ile Glu Asp Asn Gln Ile Arg Ala Ser Ser Met Leu Arg His Gly  
85 90 95

Leu Gly Ala Gln Arg Gly Arg Leu Asn Met Gln Thr Gly Ala Thr Glu  
100 105 110

Asp Asp Tyr Tyr Asp Gly Ala Trp Cys Ala Glu Asp Asp Ala Arg Thr  
115 120 125

Gln Trp Ile Glu Val Asp Thr Arg Arg Thr Thr Arg Phe Thr Gly Val  
130 135 140

Ile Thr Gln Gly Arg Asp Ser Ser Ile His Asp Asp Phe Val Thr Thr  
145 150 155 160

Phe Phe Val Gly Phe Ser Asn Asp Ser Gln Thr Trp Val Met Tyr Thr  
165 170 175

Asn Gly Tyr Glu Glu Met Thr Phe His Gly Asn Val Asp Lys Asp Thr  
180 185 190

Pro Val Leu Ser Glu Leu Pro Glu Pro Val Val Ala Arg Phe Ile Arg  
195 200 205

Ile Tyr Pro Leu Thr Trp Asn Gly Ser Leu Cys Met Arg Leu Glu Val  
210 215 220

Leu Gly Cys Ser Val Ala Pro Val Tyr Ser Tyr Tyr Ala Gln Asn Glu  
225 230 235 240

Val Val Ala Thr Asp Asp Leu Asp Phe Arg His His Ser Tyr Lys Asp  
245 250 255

Met	Arg	Gln	Leu	Met	Lys	Val	Val	Asn	Glu	Glu	Cys	Pro	Thr	Ile	Thr	260	265	270	
Arg	Thr	Tyr	Ser	Leu	Gly	Lys	Ser	Ser	Arg	Gly	Leu	Lys	Ile	Tyr	Ala	275	280	285	
Met	Glu	Ile	Ser	Asp	Asn	Pro	Gly	Glu	His	Glu	Leu	Gly	Glu	Pro	Glu	290	295	300	
Phe	Arg	Tyr	Thr	Ala	Gly	Ile	His	Gly	Asn	Glu	Val	Leu	Gly	Arg	Glu	305	310	315	320
Leu	Leu	Leu	Leu	Leu	Met	Gln	Tyr	Leu	Cys	Arg	Glu	Tyr	Arg	Asp	Gly	325	330	335	
Asn	Pro	Arg	Val	Arg	Ser	Leu	Val	Gln	Asp	Thr	Arg	Ile	His	Leu	Val	340	345	350	
Pro	Ser	Leu	Asn	Pro	Asp	Gly	Tyr	Glu	Val	Ala	Ala	Gln	Met	Gly	Ser	355	360	365	
Glu	Phe	Gly	Asn	Trp	Ala	Leu	Gly	Leu	Trp	Thr	Glu	Glu	Gly	Phe	Asp	370	375	380	
Ile	Phe	Glu	Asp	Phe	Pro	Asp	Leu	Asn	Ser	Val	Leu	Trp	Gly	Ala	Glu	385	390	395	400
Glu	Arg	Lys	Trp	Val	Pro	Tyr	Arg	Val	Pro	Asn	Asn	Asn	Leu	Pro	Ile	405	410	415	
Pro	Glu	Arg	Tyr	Leu	Ser	Pro	Asp	Ala	Thr	Val	Ser	Thr	Glu	Val	Arg	420	425	430	
Ala	Ile	Ile	Ala	Trp	Met	Glu	Lys	Asn	Pro	Phe	Val	Leu	Gly	Ala	Asn	435	440	445	
Leu	Asn	Gly	Gly	Glu	Arg	Leu	Val	Ser	Tyr	Pro	Tyr	Asp	Met	Ala	Arg	450	455	460	
Thr	Pro	Thr	Gln	Glu	Gln	Leu	Leu	Ala	Ala	Ala	Met	Ala	Ala	Ala	Arg	465	470	475	480
Gly	Glu	Asp	Glu	Asp	Glu	Val	Ser	Glu	Ala	Gln	Glu	Thr	Pro	Asp	His	485	490	495	
Ala	Ile	Phe	Arg	Trp	Leu	Ala	Ile	Ser	Phe	Ala	Ser	Ala	His	Leu	Thr	500	505	510	

Leu Thr Glu Pro Tyr Arg Gly Gly Cys Gln Ala Gln Asp Tyr Thr Gly  
515 520 525

Gly Met Gly Ile Val Asn Gly Ala Lys Trp Asn Pro Arg Thr Gly Thr  
530 535 540

Ile Asn Asp Phe Ser Tyr Leu His Thr Asn Cys Leu Glu Leu Ser Phe  
545 550 555 560

Tyr Leu Gly Cys Asp Lys Phe Pro His Glu Ser Glu Leu Pro Arg Glu  
565 570 575

Trp Glu Asn Asn Lys Glu Ala Leu Leu Thr Phe Met Glu Gln Val His  
580 585 590

Arg Gly Ile Lys Gly Val Val Thr Asp Glu Gln Gly Ile Pro Ile Ala  
595 600 605

Asn Ala Thr Ile Ser Val Ser Gly Ile Asn His Gly Val Lys Thr Ala  
610 615 620

Ser Gly Gly Asp Tyr Trp Arg Ile Leu Asn Pro Gly Glu Tyr Arg Val  
625 630 635 640

Thr Ala His Ala Glu Gly Tyr Thr Pro Ser Ala Lys Thr Cys Asn Val  
645 650 655

Asp Tyr Asp Ile Gly Ala Thr Gln Cys Asn Phe Ile Leu Ala Arg Ser  
660 665 670

Asn Trp Lys Arg Ile Arg Glu Ile Met Ala Met Asn Gly Asn Arg Pro  
675 680 685

Ile Pro His Ile Asp Pro Ser Arg Pro Met Thr Pro Gln Gln Arg Arg  
690 695 700

Leu Gln Gln Arg Arg Leu Gln His Arg Leu Arg Leu Arg Ala Gln Met  
705 710 715 720

Arg Leu Arg Arg Leu Asn Ala Thr Thr Thr Leu Gly Pro His Thr Val  
725 730 735

Pro Pro Thr Leu Pro Pro Ala Pro Ala Thr Thr Leu Ser Thr Thr Ile  
740 745 750

Glu Pro Trp Gly Leu Ile Pro Pro Thr Thr Ala Gly Trp Glu Glu Ser  
755 760 765

Glu Thr Glu Thr Tyr Thr Glu Val Val Thr Glu Phe Gly Thr Glu Val  
770 775 780

Glu Pro Glu Phe Gly Thr Lys Val Glu Pro Glu Phe Glu Thr Gln Leu  
785 790 795 800

Glu Pro Glu Phe Glu Thr Gln Leu Glu Pro Glu Phe Glu Glu Glu Glu  
805 810 815

Glu Glu Glu Lys Glu Glu Glu Ile Ala Thr Gly Gln Ala Phe Pro Phe  
820 825 830

Thr Thr Val Glu Thr Tyr Thr Val Asn Phe Gly Asp Phe  
835 840 845

<210> 12

<211> 764

<212> PRT

<213> Mus musculus

<400> 12

Met Ala Arg Leu Gly Thr Ala Cys Pro Ala Leu Ala Leu Ala Leu Ala  
1 5 10 15

Leu Val Ala Val Ala Leu Ala Gly Val Arg Ala Gln Gly Ala Ala Phe  
20 25 30

Glu Glu Pro Asp Tyr Tyr Ser Gln Glu Leu Trp Arg Arg Gly Arg Tyr  
35 40 45

Tyr Gly His Pro Glu Pro Glu Pro Glu Pro Glu Leu Phe Ser Pro Ser  
50 55 60

Met His Glu Asp Leu Arg Val Glu Glu Gln Glu Gln Gln Arg Pro His  
65 70 75 80

Gln Gln Gly His Arg Thr Pro Lys Lys Ala Ile Lys Pro Lys Lys Ala  
85 90 95

Pro Lys Arg Glu Lys Leu Val Ala Glu Thr Pro Pro Pro Gly Lys Asn  
100 105 110

Ser Asn Arg Lys Gly Arg Arg Ser Lys Asn Leu Glu Lys Ala Ala Ser  
115 120 125

Asp Asp His Gly Val Pro Val Ala His Glu Asp Val Arg Glu Ser Cys

130		135		140	
Pro Pro Leu Gly Leu Glu Thr Leu Lys Ile Thr Asp Phe Gln Leu His					
145		150		155	160
Ala Ser Thr Ser Lys Arg Tyr Gly Leu Gly Ala His Arg Gly Arg Leu					
	165		170		175
Asn Ile Gln Ala Gly Ile Asn Glu Asn Asp Phe Tyr Asp Gly Ala Trp					
	180		185		190
Cys Ala Gly Arg Asn Asp Leu His Gln Trp Ile Glu Val Asp Ala Arg					
	195		200		205
Arg Leu Thr Lys Phe Thr Gly Val Ile Thr Gln Gly Arg Asn Ser Leu					
	210		215		220
Trp Leu Ser Asp Trp Val Thr Ser Tyr Lys Val Met Val Ser Asn Asp					
	225		230		235
Ser His Thr Trp Val Thr Val Lys Asn Gly Ser Gly Asp Met Ile Phe					
	245		250		255
Glu Gly Asn Ser Glu Lys Glu Ile Pro Val Leu Asn Glu Leu Pro Val					
	260		265		270
Pro Met Val Ala Arg Val Ile Arg Ile Asn Pro Gln Ser Trp Phe Asp					
	275		280		285
Asn Gly Ser Ile Cys Met Arg Met Glu Ile Leu Gly Cys Pro Leu Pro					
	290		295		300
Asp Pro Asn Asn Tyr Tyr His Arg Arg Asn Glu Met Thr Thr Thr Asp					
	305		310		315
Asp Leu Asp Phe Lys His His Asn Tyr Lys Glu Met Arg Gln Leu Met					
	325		330		335
Lys Val Val Asn Glu Met Cys Pro Asn Ile Thr Arg Ile Tyr Asn Ile					
	340		345		350
Gly Lys Ser His Gln Gly Leu Lys Leu Tyr Ala Val Glu Ile Ser Asp					
	355		360		365
His Pro Gly Glu His Glu Val Gly Glu Pro Glu Phe His Tyr Ile Ala					
	370		375		380
Gly Ala His Gly Asn Glu Val Leu Gly Arg Glu Leu Leu Leu Leu Leu					



385		390		395		400
Leu His Phe Leu Cys Gln Glu Tyr Ser Ala Gln Asn Ala Arg Ile Val						
	405		410		415	
Arg Leu Val Glu Glu Thr Arg Ile His Ile Leu Pro Ser Leu Asn Pro						
	420		425		430	
Asp Val Tyr Glu Lys Ala Tyr Glu Gly Gly Ser Glu Leu Gly Gly Trp						
	435		440		445	
Ser Leu Gly Arg Trp Thr His Asp Gly Ile Asp Ile Asn Asn Asn Phe						
	450		455		460	
Pro Asp Leu Asn Ser Leu Leu Trp Glu Ala Glu Asp Gln Gln Asn Ala						
465		470		475		480
Pro Arg Lys Val Pro Asn His Tyr Ile Ala Ile Pro Glu Trp Phe Leu						
	485		490		495	
Ser Glu Asn Ala Thr Val Ala Thr Glu Thr Arg Ala Val Ile Ala Trp						
	500		505		510	
Met Glu Lys Ile Pro Phe Val Leu Gly Gly Asn Leu Gln Gly Gly Glu						
	515		520		525	
Leu Val Val Ala Tyr Pro Tyr Asp Met Val Arg Ser Leu Trp Lys Thr						
	530		535		540	
Gln Glu His Thr Pro Thr Pro Asp Asp His Val Phe Arg Trp Leu Ala						
545		550		555		560
Tyr Ser Tyr Ala Ser Thr His Arg Leu Met Thr Asp Ala Arg Arg Arg						
	565		570		575	
Val Cys His Thr Glu Asp Phe Gln Lys Glu Glu Gly Thr Val Asn Gly						
	580		585		590	
Ala Ser Trp His Thr Val Ala Gly Ser Leu Asn Asp Phe Ser Tyr Leu						
	595		600		605	
Gly Thr Asn Cys Phe Glu Leu Ser Ile Tyr Val Gly Cys Asp Lys Tyr						
	610		615		620	
Pro His Glu Ser Glu Leu Pro Glu Glu Trp Glu Asn Asn Arg Glu Ser						
625		630		635		640
Leu Ile Val Phe Met Glu Gln Val His Arg Gly Ile Lys Gly Ile Val						

	645		650		655
Arg Asp Leu Gln Gly Lys Gly Ile Ser Asn Ala Val Ile Ser Val Glu					
	660		665		670
Gly Val Asn His Asp Ile Arg Thr Ala Ser Asp Gly Asp Tyr Trp Arg					
	675		680		685
Leu Leu Asn Pro Gly Glu Tyr Val Val Thr Ala Lys Ala Glu Gly Phe					
	690		695		700
Ile Thr Ser Thr Lys Asn Cys Met Val Gly Tyr Asp Met Gly Ala Thr					
	705		710		715
Arg Cys Asp Phe Thr Leu Thr Lys Thr Asn Leu Ala Arg Ile Arg Glu					
	725		730		735
Ile Met Glu Thr Phe Gly Lys Gln Pro Val Ser Leu Pro Ser Arg Arg					
	740		745		750
Leu Lys Leu Arg Gly Arg Lys Arg Arg Gln Arg Gly					
	755		760		

<210> 13  
 <211> 722  
 <212> PRT  
 <213> Mus musculus

<400> 13  
 Met Trp Gly Leu Leu Leu Ala Val Thr Ala Phe Ala Pro Ser Val Gly  
 1 5 10 15

Leu Gly Leu Gly Ala Pro Ser Ala Ser Val Pro Gly Leu Ala Pro Gly  
 20 25 30

Ser Thr Leu Ala Pro His Ser Ser Val Ala Gln Pro Ser Thr Lys Ala  
 35 40 45

Asn Glu Thr Ser Glu Arg His Val Arg Leu Arg Val Ile Lys Lys Lys  
 50 55 60

Lys Ile Val Val Lys Lys Arg Lys Lys Leu Arg His Pro Gly Pro Leu  
 65 70 75 80

Gly Thr Ala Arg Pro Val Val Pro Thr His Pro Ala Lys Thr Leu Thr  
 85 90 95

Leu Pro Glu Lys Gln Glu Pro Gly Cys Pro Pro Leu Gly Leu Glu Ser  
 100 105 110

Leu Arg Val Ser Asp Ser Gln Leu Glu Ala Ser Ser Ser Gln Ser Phe  
 115 120 125

Gly Leu Gly Ala His Arg Gly Arg Leu Asn Ile Gln Ser Gly Leu Glu  
 130 135 140

Asp Gly Asp Leu Tyr Asp Gly Ala Trp Cys Ala Glu Gln Gln Asp Thr  
 145 150 155 160

Glu Pro Trp Leu Gln Val Asp Ala Lys Asn Pro Val Arg Phe Ala Gly  
 165 170 175

Ile Val Thr Gln Gly Arg Asn Ser Val Trp Arg Tyr Asp Trp Val Thr  
 180 185 190

Ser Phe Lys Val Gln Phe Ser Asn Asp Ser Gln Thr Trp Trp Lys Ser  
 195 200 205

Arg Asn Ser Thr Gly Met Asp Ile Val Phe Pro Ala Asn Ser Asp Ala  
 210 215 220

Glu Thr Pro Val Leu Asn Leu Leu Pro Glu Pro Gln Val Ala Arg Phe  
 225 230 235 240

Ile Arg Leu Leu Pro Gln Thr Trp Phe Gln Gly Gly Val Pro Cys Leu  
 245 250 255

Arg Ala Glu Ile Leu Ala Cys Pro Val Ser Asp Pro Asn Asp Leu Phe  
 260 265 270

Pro Glu Ala His Thr Leu Gly Ser Ser Asn Ser Leu Asp Phe Arg His  
 275 280 285

His Asn Tyr Lys Ala Met Arg Lys Leu Met Lys Gln Val Asn Glu Gln  
 290 295 300

Cys Pro Asn Ile Thr Arg Ile Tyr Ser Ile Gly Lys Ser His Gln Gly  
 305 310 315 320

Leu Lys Leu Tyr Val Met Glu Met Ser Asp His Pro Gly Glu His Glu  
 325 330 335

Leu Gly Glu Pro Glu Val Arg Tyr Val Ala Gly Met His Gly Asn Glu  
 340 345 350

Ala	Leu	Gly	Arg	Glu	Leu	Leu	Leu	Leu	Leu	Met	Gln	Phe	Leu	Cys	His
	355						360					365			
Glu	Phe	Leu	Arg	Gly	Asp	Pro	Arg	Val	Thr	Arg	Leu	Leu	Thr	Glu	Thr
	370					375					380				
Arg	Ile	His	Leu	Leu	Pro	Ser	Met	Asn	Pro	Asp	Gly	Tyr	Glu	Thr	Ala
385					390					395					400
Tyr	His	Arg	Gly	Ser	Glu	Leu	Val	Gly	Trp	Ala	Glu	Gly	Arg	Trp	Thr
			405					410					415		
His	Gln	Gly	Ile	Asp	Leu	Asn	His	Asn	Phe	Ala	Asp	Leu	Asn	Thr	Gln
		420					425					430			
Leu	Trp	Tyr	Ala	Glu	Asp	Asp	Gly	Leu	Val	Pro	Asp	Thr	Val	Pro	Asn
	435						440					445			
His	His	Leu	Pro	Leu	Pro	Thr	Tyr	Tyr	Thr	Leu	Pro	Asn	Ala	Thr	Val
	450					455					460				
Ala	Pro	Glu	Thr	Trp	Ala	Val	Ile	Lys	Trp	Met	Lys	Arg	Ile	Pro	Phe
465					470					475					480
Val	Leu	Ser	Ala	Asn	Leu	His	Gly	Gly	Glu	Leu	Val	Val	Ser	Tyr	Pro
			485					490					495		
Phe	Asp	Met	Thr	Arg	Thr	Pro	Trp	Ala	Ala	Arg	Glu	Leu	Thr	Pro	Thr
		500						505					510		
Pro	Asp	Asp	Ala	Val	Phe	Arg	Trp	Leu	Ser	Thr	Val	Tyr	Ala	Gly	Thr
	515						520					525			
Asn	Arg	Ala	Met	Gln	Asp	Thr	Asp	Arg	Arg	Pro	Cys	His	Ser	Gln	Asp
	530					535					540				
Phe	Ser	Leu	His	Gly	Asn	Val	Ile	Asn	Gly	Ala	Asp	Trp	His	Thr	Val
545					550				555					560	
Pro	Gly	Ser	Met	Asn	Asp	Phe	Ser	Tyr	Leu	His	Thr	Asn	Cys	Phe	Glu
			565					570					575		
Val	Thr	Val	Glu	Leu	Ser	Cys	Asp	Lys	Phe	Pro	His	Glu	Lys	Glu	Leu
		580						585					590		
Pro	Gln	Glu	Trp	Glu	Asn	Asn	Lys	Asp	Ala	Leu	Leu	Thr	Tyr	Leu	Glu
	595						600					605			

Gln Val Arg Met Gly Ile Thr Gly Val Val Arg Asp Lys Asp Thr Glu  
610 615 620

Leu Gly Ile Ala Asp Ala Val Ile Ala Val Glu Gly Ile Asn His Asp  
625 630 635 640

Val Thr Thr Ala Trp Gly Gly Asp Tyr Trp Arg Leu Leu Thr Pro Gly  
645 650 655

Asp Tyr Val Val Thr Ala Ser Ala Glu Gly Tyr His Thr Val Arg Gln  
660 665 670

His Cys Gln Val Thr Phe Glu Glu Gly Pro Val Pro Cys Asn Phe Leu  
675 680 685

Leu Thr Lys Thr Pro Lys Glu Arg Leu Arg Glu Leu Leu Ala Thr Arg  
690 695 700

Gly Lys Leu Pro Pro Asp Leu Arg Arg Lys Leu Glu Arg Leu Arg Gly  
705 710 715 720

Gln Lys

<210> 14

<211> 40

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 14

ctcgtcagat ctgcgcccag gaactcggtg ctgggcctcg

40

<210> 15

<211> 37

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 15

ctcgtectcg agatccttct gtccccttag ccgctcc

37

<210> 16

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 16

agccggcttg aggcattcag c

21

<210> 17

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 17

gttgatgcc tcaagccggc t

21

<210> 18

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 18

ccagaaactc cagtgtgaa c

21

<210> 19

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 19

gttcagcact ggagtttctg g

21

<210> 20

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 20

caagcctggg gagcatgagc tg

22

<210> 21

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 21

cagctcatgc tccccaggt tg

22

<210> 22

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 22

caggacgatg ggaaggtgcc c

21

<210> 23

<211> 21

<212> DNA  
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 23  
gggcaccttc ccacgtcct g

21

<210> 24  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 24  
agcatgaatg aattcagcta c

21

<210> 25  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 25  
gtagctgaag tcattcatgc t

21

<210> 26  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 26  
gagcttgga ttgctgacgc t

21



<210> 27  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 27  
gggtcagcaa tcccaagctc 20

<210> 28  
<211> 30  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 28  
ctcgtcctcg agggtaagcc tatccctaac 30

<210> 29  
<211> 31  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 29  
ctcgtcgggc cctgatcag cgggtttaaa c 31

<210> 30  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Chemically

Synthesized

<400> 30  
gtctggagtc cctgcgagtt t 21

<210> 31  
<211> 19  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 31  
cggtgtggtc caagaccaa 19

<210> 32  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 32  
cttgaggcat ccagcagcca gtcc 24

<210> 33  
<211> 23  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 33  
gagtcctgc gagtttcaga tag 23

<210> 34  
<211> 20  
<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 34

gtcctcggtg tggccaaga

20

<210> 35

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 35

tgaggcatcc agcagccagt ccttt

25

<210> 36

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 36

gtgctgctgc tctacaataa cca

23

<210> 37

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 37

gtttctgcag ctgggcat

19

<210> 38  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 38  
tggaccggtg cgccttcgat 20

<210> 39  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 39  
gacgtgggat gcacacagct c 21

<210> 40  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Chemically  
Synthesized

<400> 40  
gttgcaactgc tctgggggtca 20

<210> 41  
<211> 67  
<212> PRT  
<213> Homo sapiens

<400> 41  
Pro Asp Met Val Thr Ala Ser Ala Glu Gly Tyr His Ser Val Thr Arg  
1 5 10 15

Asn	Cys	Arg	Val	Thr	Phe	Glu	Glu	Gly	Pro	Phe	Pro	Cys	Asn	Phe	Val
			20					25					30		

Leu	Thr	Lys	Thr	Pro	Lys	Gln	Arg	Leu	Arg	Glu	Leu	Leu	Ala	Ala	Gly
		35					40				45				

Ala	Lys	Val	Pro	Pro	Asp	Leu	Arg	Arg	Arg	Leu	Glu	Arg	Leu	Arg	Gly
	50					55					60				

Gln	Lys	Asp
65		

<210> 42  
 <211> 69  
 <212> PRT  
 <213> Homo sapiens

<400> 42															
Pro	Gly	Asp	Tyr	Met	Val	Thr	Ala	Ser	Ala	Glu	Gly	Tyr	His	Ser	Val
1				5					10					15	

Thr	Arg	Asn	Cys	Arg	Val	Thr	Phe	Glu	Glu	Gly	Pro	Phe	Pro	Cys	Asn
			20					25					30		

Phe	Val	Leu	Thr	Lys	Thr	Pro	Lys	Gln	Arg	Leu	Arg	Glu	Leu	Leu	Ala
		35						40				45			

Ala	Gly	Ala	Lys	Val	Pro	Pro	Asp	Leu	Arg	Arg	Arg	Leu	Glu	Arg	Leu
	50						55					60			

Arg	Gly	Gln	Lys	Asp
65				

<210> 43  
 <211> 192  
 <212> PRT  
 <213> Homo sapiens

<400> 43															
Met	Trp	Gly	Leu	Leu	Leu	Ala	Leu	Ala	Ala	Phe	Ala	Pro	Ala	Val	Gly
1				5					10					15	

Pro	Ala	Leu	Gly	Ala	Pro	Arg	Asn	Ser	Val	Leu	Gly	Leu	Ala	Gln	Pro
			20					25					30		

Gly Thr Thr Lys Val Pro Gly Ser Thr Pro Ala Leu His Ser Ser Pro  
 35 40 45

Ala Gln Pro Pro Ala Glu Thr Ala Asn Gly Thr Ser Glu Gln His Val  
 50 55 60

Arg Ile Arg Val Ile Lys Lys Lys Lys Val Ile Met Lys Lys Arg Lys  
 65 70 75 80

Lys Leu Thr Leu Thr Arg Pro Thr Pro Leu Val Thr Ala Gly Pro Leu  
 85 90 95

Val Thr Pro Thr Pro Ala Gly Thr Leu Asp Pro Ala Glu Lys Gln Glu  
 100 105 110

Thr Gly Cys Pro Pro Leu Gly Leu Glu Ser Leu Arg Val Ser Asp Ser  
 115 120 125

Arg Leu Glu Ala Ser Ser Ser Gln Ser Phe Gly Leu Gly Pro His Arg  
 130 135 140

Gly Arg Leu Asn Ile Gln Ser Gly Leu Glu Asp Gly Asp Leu Tyr Asp  
 145 150 155 160

Gly Ala Trp Cys Ala Glu Glu Gln Asp Ala Asp Pro Trp Phe Gln Val  
 165 170 175

Asp Ala Gly His Pro Thr Arg Phe Ser Gly Val Ile Thr Gln Gly Arg  
 180 185 190

<210> 44

<211> 193

<212> PRT

<213> Homo sapiens

<400> 44

Met Trp Gly Leu Leu Leu Ala Leu Ala Ala Phe Ala Pro Ala Val Gly  
 1 5 10 15

Pro Ala Leu Gly Ala Pro Arg Asn Ser Val Leu Gly Leu Ala Gln Pro  
 20 25 30

Gly Thr Thr Lys Val Pro Gly Ser Thr Pro Ala Leu His Ser Ser Pro  
 35 40 45

Ala Gln Pro Pro Ala Glu Thr Ala Asn Gly Thr Ser Glu Gln His Val  
50 55 60

Arg Ile Arg Val Ile Lys Lys Lys Lys Val Ile Met Lys Lys Arg Lys  
65 70 75 80

Lys Leu Thr Leu Thr Arg Pro Thr Pro Leu Val Thr Ala Gly Pro Leu  
85 90 95

Val Thr Pro Thr Pro Ala Gly Thr Leu Asp Pro Ala Glu Lys Gln Glu  
100 105 110

Thr Gly Cys Pro Pro Leu Gly Leu Glu Ser Leu Arg Val Ser Asp Ser  
115 120 125

Arg Leu Glu Ala Ser Ser Ser Gln Ser Phe Gly Leu Gly Pro His Arg  
130 135 140

Gly Arg Leu Asn Ile Gln Ser Gly Leu Glu Asp Gly Asp Leu Tyr Asp  
145 150 155 160

Gly Ala Trp Cys Ala Glu Glu Gln Asp Ala Asp Pro Trp Phe Gln Val  
165 170 175

Asp Ala Gly His Pro Thr Arg Phe Ser Gly Val Ile Thr Gln Gly Arg  
180 185 190

Asn

<210> 45

<211> 510

<212> PRT

<213> Homo sapiens

<400> 45

Met Trp Gly Leu Leu Leu Ala Leu Ala Ala Phe Ala Pro Ala Val Gly  
1 5 10 15

Pro Ala Leu Gly Ala Pro Arg Asn Ser Val Leu Gly Leu Ala Gln Pro  
20 25 30

Gly Thr Thr Lys Val Pro Gly Ser Thr Pro Ala Leu His Ser Ser Pro  
35 40 45

Ala Gln Pro Pro Ala Glu Thr Ala Asn Gly Thr Ser Glu Gln His Val

50	55	60
Arg Ile Arg Val Ile Lys Lys Lys Lys Val Ile Met Lys Lys Arg Lys		
65	70	75 80
Lys Leu Thr Leu Thr Arg Pro Thr Pro Leu Val Thr Ala Gly Pro Leu		
85	90	95
Val Thr Pro Thr Pro Ala Gly Thr Leu Asp Pro Ala Glu Lys Gln Glu		
100	105	110
Thr Gly Cys Pro Pro Leu Gly Leu Glu Ser Leu Arg Val Ser Asp Ser		
115	120	125
Arg Leu Glu Ala Ser Ser Ser Gln Ser Phe Gly Leu Gly Pro His Arg		
130	135	140
Gly Arg Leu Asn Ile Gln Ser Gly Leu Glu Asp Gly Asp Leu Tyr Asp		
145	150	155 160
Gly Ala Trp Cys Ala Glu Glu Gln Asp Ala Asp Pro Trp Phe Gln Val		
165	170	175
Asp Ala Gly His Pro Thr Arg Phe Ser Gly Val Ile Thr Gln Gly Arg		
180	185	190
Asn Ser Val Trp Arg Tyr Asp Trp Val Thr Ser Tyr Lys Val Gln Phe		
195	200	205
Ser Asn Asp Ser Arg Thr Trp Trp Gly Ser Arg Asn His Ser Ser Gly		
210	215	220
Met Asp Ala Val Phe Pro Ala Asn Ser Asp Pro Glu Thr Pro Val Leu		
225	230	235 240
Asn Leu Leu Pro Glu Pro Gln Val Ala Arg Phe Ile Arg Leu Leu Pro		
245	250	255
Gln Thr Trp Leu Gln Gly Gly Ala Pro Cys Leu Arg Ala Glu Ile Leu		
260	265	270
Ala Cys Pro Val Ser Asp Pro Asn Asp Leu Phe Leu Glu Ala Pro Ala		
275	280	285
Ser Gly Ser Ser Asp Pro Leu Asp Phe Gln His His Asn Tyr Lys Ala		
290	295	300
Met Arg Lys Leu Met Lys Gln Val Gln Glu Gln Cys Pro Asn Ile Thr		





Arg Asn Cys Arg Val Thr Phe Glu Glu Gly Pro Phe Pro Cys Asn Phe  
 20 25 30

Val Leu Thr Lys Thr Pro Lys Gln Arg Leu Arg Glu Leu Leu Ala Ala  
 35 40 45

Gly Ala Lys Val Pro Pro Asp Leu Arg Arg Arg Leu Glu Arg Leu Arg  
 50 55 60

Gly Gln Lys Asp  
 65

<210> 47

<211> 193

<212> PRT

<213> Homo sapiens

<400> 47

Met Trp Gly Leu Leu Leu Ala Leu Ala Ala Phe Ala Pro Ala Val Gly  
 1 5 10 15

Pro Ala Leu Gly Ala Pro Arg Asn Ser Val Leu Gly Leu Ala Gln Pro  
 20 25 30

Gly Thr Thr Lys Val Pro Gly Ser Thr Pro Ala Leu His Ser Ser Pro  
 35 40 45

Ala Gln Pro Pro Ala Glu Thr Ala Asn Gly Thr Ser Glu Gln His Val  
 50 55 60

Arg Ile Arg Val Ile Lys Lys Lys Lys Val Ile Met Lys Lys Arg Lys  
 65 70 75 80

Lys Leu Thr Leu Thr Arg Pro Thr Pro Leu Val Thr Ala Gly Pro Leu  
 85 90 95

Val Thr Pro Thr Pro Ala Gly Thr Leu Asp Pro Ala Glu Lys Gln Glu  
 100 105 110

Thr Gly Cys Pro Pro Leu Gly Leu Glu Ser Leu Arg Val Ser Asp Ser  
 115 120 125

Arg Leu Glu Ala Ser Ser Ser Gln Ser Phe Gly Leu Gly Pro His Arg  
 130 135 140

Gly Arg Leu Asn Ile Gln Ser Gly Leu Glu Asp Gly Asp Leu Tyr Asp  
 145 150 155 160

Gly Ala Trp Cys Ala Glu Glu Gln Asp Ala Asp Pro Trp Phe Gln Val  
165 170 175

Asp Ala Gly His Pro Thr Arg Phe Ser Gly Val Ile Thr Gln Gly Arg  
180 185 190

Asp